

## SEQUENCE LISTING

Merman, John L Rastelli, Luca Shimkets, Richard A

<120> Novel Proteins and Nucleic Acids Encoding Same and Antibodies Directed Against these Proteins

<130> 21402-138

<140> 09/970,944

<141> 2001-10-04

<150> 60/237,862

<151> 2000-10-04

<160> 62

<170> PatentIn Ver. 2.1

<210> 1

<211> 2881

<212> DNA

<213> Homo sapiens

<400> 1

agctggggct ccgggctgag gcgctaaagc cgccctcccg cccgcggggc cccgcgcccg 60 gcccgcccgc ctgcccgccc gcggccatgg ccgtccggcc cggcctgtgg ccagcgctcc 120 tgggcatagt cctcgccgct tggctccgcg gctcgggtgc ccagcagagt gccaccgtgg 180 ccaacccagt gcctggtgcc aacccggacc tgcttcccca cttcctggtg gagcccgagg 240 atgtgtacat cgtcaagaac aagccagtgc tgcttgtgtg caaggccgtg cccgccacgc 300 agatettett caagtgeaac ggggagtggg tgegeeaggt ggaceaegtg ategagegea 360 gcacagacgg gagcagtggt gagccgacca tggaggtccg cattaatgtc tcaaggcagc 420 aggtcgagaa ggtgttcggg ctggaggaat actggtgcca gtgcgtggca tggagctcct 480 cgggcaccac caagagtcag aaggcctaca tccgcatagc cagattgcgc aagaacttcg 540 agcaggagcc gctggccaag gaggtgtccc tggagcaggg catcgtgctg ccctgccgtc 600 caccggaggg catccctcca gccgaggtgg agtggctccg gaacgaggac ctggtggacc 660 cgtccctgga ccccaatgta tacatcacgc gggagcacag cctggtggtg cgacaggccc 720 gccttgctga cacggccaac tacacctgcg tggccaagaa catcgtggca cgtcgccgca 780 gegeeteege tgetgteate gtetaegtga aeggtgggtg gtegaegtgg aeegagtggt 840 cegtetgeag egecagetgt gggegegget ggcagaaaeg gageeggage tgeaceaace 900 cggcgcctct caacgggggc gctttctgtg aggggcagaa tgtccatgac cgcaccgtct 960 cetetetget tgtetetgtg gacggcaget ggagcccgtg gagcaagtgg tcggcetgtg 1020 ggctggactg cacccactgg cggagccgtg agtgctctga cccagcaccc cgcaacggag 1080 gggaggagtg ccagggcact gacctggaca cccgcaactg taccagtgac ctctgtgtac 1140 acagtgette tggccetgag gaegtggeee tetatgtggg ceteategee gtggeegtet 1200 gcctggtcct gctgctgctt gtcctcatcc tcgtttattg ccggaagaag gaggggctgg 1260 actcagatgt ggctgactcg tccattctca cctcaggctt ccagcccgtc agcatcaagc 1320 ccagcaaagc agacaaccc catctgctca ccatccagc ggacctcagc accaccacca 1380 cctaccaggg cagtetetgt ecceggeagg atgggeecag ecceaagtte cageteacca 1440 atgggcacet geteageece etgggtggeg geegecacae actgeaceae ageteteeca 1500 cetetgagge egaggagtte gtetecegee tetecaceca gaactaette egeteeetge 1560 cccgaggcac cagcaacatg acctatggga ccttcaactt cctcgggggc cggctgatga 1620 tecetaatae aggtateage etecteatee eeccagatge cataceeega gggaagatet 1680 atgagateta ceteaegetg cacaageegg aagaegtgag gttgeeecta getggetgte 1740 agaccetget gagteceate gttagetgtg gaccecetgg egtectgete acceggecag 1800

```
teateetgge tatggaceae tgtggggage eeageeetga eagetggage etgegeetea 1860
 aaaagcagtc gtgcgaggc agctgggagc aggatgtgct gcacctgggc gaggaggcgc 1920
 cctcccacct ctactactgc cagctggagg ccagtgcctg ctacgtcttc accgagcagc 1980
tgggccgctt tgccctggtg ggagaggccc tcagcgtggc tgccgccaag cgcctcaagc 2040
tgcttctgtt tgcgccggtg gcctgcacct ccctcgagta caacatccgg gtctactgcc 2100
tgcatgacac ccacgatgca ctcaaggagg tggtgcagct ggagaagcag ctggggggac 2160
agctgateca ggagecaegg gteetgeact teaaggacag ttaccaeaac etgegeetat 2220
ccatccacga tgtgcccagc tccctgtgga agagtaagct ccttgtcagc taccaggaga 2280
teceetttta teacatetgg aatggeacge ageggtaett geactgeace tteaceetgg 2340
agegtgteag ecceageact agtgacetgg ectgeaaget gtgggtgtgg caggtggagg 2400
gegacgggca gagetteage ateaacttea acateaceaa ggacacaagg tttgetgage 2460
tgctggctct ggagagtgaa gcgggggtcc cagccctggt gggccccagt gccttcaaga 2520
teceetteet catteggeag aagataattt ceageetgga eecaeeetgt aggegggtg 2580
ccgactggcg gactctggcc cagaaactcc acctggacag ccatctcagc ttctttgcct 2640
ccaagcccag ccccacagcc atgatcctca acctgtggga ggcgcggcac ttccccaacg 2700
gcaacctcag ccagctggct gcagcagtgg ctggactggg ccagccagac gctggcctct 2760
tcacagtgtc ggaggctgag tgctgaggcc ggccaggccc gacacctaca ctctcaccag 2820
ctttggcacc caccaaggac aggcagaagc cggacagggg cccttcccca caccggggag 2880
<210> 2
<211> 899
<212> PRT
<213> Homo sapiens
<400> 2
Met Ala Val Arg Pro Gly Leu Trp Pro Ala Leu Leu Gly Ile Val Leu
Ala Ala Trp Leu Arg Gly Ser Gly Ala Gln Gln Ser Ala Thr Val Ala
```

2.0

Asn Pro Val Pro Gly Ala Asn Pro Asp Leu Leu Pro His Phe Leu Val

Glu Pro Glu Asp Val Tyr Ile Val Lys Asn Lys Pro Val Leu Leu Val 55

Cys Lys Ala Val Pro Ala Thr Gln Ile Phe Phe Lys Cys Asn Gly Glu 65 70

Trp Val Arg Gln Val Asp His Val Ile Glu Arg Ser Thr Asp Gly Ser

Ser Gly Glu Pro Thr Met Glu Val Arg Ile Asn Val Ser Arg Gln Gln 100 105

Val Glu Lys Val Phe Gly Leu Glu Glu Tyr Trp Cys Gln Cys Val Ala 115

Trp Ser Ser Ser Gly Thr Thr Lys Ser Gln Lys Ala Tyr Ile Arg Ile 135

Ala Arg Leu Arg Lys Asn Phe Glu Gln Glu Pro Leu Ala Lys Glu Val 145 150 160

- Ser Leu Glu Gln Gly Ile Val Leu Pro Cys Arg Pro Pro Glu Gly Ile 165 170 175
- Pro Pro Ala Glu Val Glu Trp Leu Arg Asn Glu Asp Leu Val Asp Pro 180 185 190
- Ser Leu Asp Pro Asn Val Tyr Ile Thr Arg Glu His Ser Leu Val Val 195 200 205
- Arg Gln Ala Arg Leu Ala Asp Thr Ala Asn Tyr Thr Cys Val Ala Lys 210 215 220
- Asn Ile Val Ala Arg Arg Arg Ser Ala Ser Ala Ala Val Ile Val Tyr 225 230 235 240
- Val Asn Gly Gly Trp Ser Thr Trp Thr Glu Trp Ser Val Cys Ser Ala 245 250 255
- Ser Cys Gly Arg Gly Trp Gln Lys Arg Ser Arg Ser Cys Thr Asn Pro 260 265 270
- Ala Pro Leu Asn Gly Gly Ala Phe Cys Glu Gly Gln Asn Val His Asp 275 280 285
- Arg Thr Val Ser Ser Leu Leu Val Ser Val Asp Gly Ser Trp Ser Pro 290 295 300
- Trp Ser Lys Trp Ser Ala Cys Gly Leu Asp Cys Thr His Trp Arg Ser 305 310 310 315 320
- Arg Glu Cys Ser Asp Pro Ala Pro Arg Asn Gly Gly Glu Glu Cys Gln 325 330 335
- Gly Thr Asp Leu Asp Thr Arg Asn Cys Thr Ser Asp Leu Cys Val His 340 345 350
- Ser Ala Ser Gly Pro Glu Asp Val Ala Leu Tyr Val Gly Leu Ile Ala 355 360 365
- Val Ala Val Cys Leu Val Leu Leu Leu Val Leu Ile Leu Val Tyr 370 375 380
- Cys Arg Lys Lys Glu Gly Leu Asp Ser Asp Val Ala Asp Ser Ser Ile 385 390 395 400
- Leu Thr Ser Gly Phe Gln Pro Val Ser Ile Lys Pro Ser Lys Ala Asp 405 410 415
- Asn Pro His Leu Leu Thr Ile Gln Pro Asp Leu Ser Thr Thr Thr 420 425 430
- Tyr Gln Gly Ser Leu Cys Pro Arg Gln Asp Gly Pro Ser Pro Lys Phe 435
- Gln Leu Thr Asn Gly His Leu Leu Ser Pro Leu Gly Gly Gly Arg His 450 455 460

- Thr Leu His His Ser Ser Pro Thr Ser Glu Ala Glu Glu Phe Val Ser 465 470 475 480
- Arg Leu Ser Thr Gln Asn Tyr Phe Arg Ser Leu Pro Arg Gly Thr Ser 485 490 495
- Asn Met Thr Tyr Gly Thr Phe Asn Phe Leu Gly Gly Arg Leu Met Ile 500 505 510
- Pro Asn Thr Gly Ile Ser Leu Leu Ile Pro Pro Asp Ala Ile Pro Arg 515 520 525
- Gly Lys Ile Tyr Glu Ile Tyr Leu Thr Leu His Lys Pro Glu Asp Val 530 535 540
- Arg Leu Pro Leu Ala Gly Cys Gln Thr Leu Leu Ser Pro Ile Val Ser 545 550 555 560
- Cys Gly Pro Pro Gly Val Leu Leu Thr Arg Pro Val Ile Leu Ala Met 565 570 575
- Asp His Cys Gly Glu Pro Ser Pro Asp Ser Trp Ser Leu Arg Leu Lys
  580 585 590
- Lys Gln Ser Cys Glu Gly Ser Trp Glu Gln Asp Val Leu His Leu Gly 595 600 605
- Glu Glu Ala Pro Ser His Leu Tyr Tyr Cys Gln Leu Glu Ala Ser Ala 610 615 620
- Cys Tyr Val Phe Thr Glu Gln Leu Gly Arg Phe Ala Leu Val Gly Glu 625 630 635 640
- Ala Leu Ser Val Ala Ala Ala Lys Arg Leu Lys Leu Leu Leu Phe Ala 645 650 655
- Pro Val Ala Cys Thr Ser Leu Glu Tyr Asn Ile Arg Val Tyr Cys Leu 660 665 670
- His Asp Thr His Asp Ala Leu Lys Glu Val Val Gln Leu Glu Lys Gln 675 680 685
- Leu Gly Gly Gln Leu Ile Gln Glu Pro Arg Val Leu His Phe Lys Asp
  690 695 700
- Ser Tyr His Asn Leu Arg Leu Ser Ile His Asp Val Pro Ser Ser Leu 705 710 715 720
- Trp Lys Ser Lys Leu Leu Val Ser Tyr Gln Glu Ile Pro Phe Tyr His 725 730 735
- Ile Trp Asn Gly Thr Gln Arg Tyr Leu His Cys Thr Phe Thr Leu Glu 740 745 750
- Arg Val Ser Pro Ser Thr Ser Asp Leu Ala Cys Lys Leu Trp Val Trp
  755 760 765

```
Gln Val Glu Gly Asp Gly Gln Ser Phe Ser Ile Asn Phe Asn Ile Thr
     770
Lys Asp Thr Arg Phe Ala Glu Leu Leu Ala Leu Glu Ser Glu Ala Gly
785
                     790
                                         795
                                                              800
Val Pro Ala Leu Val Gly Pro Ser Ala Phe Lys Ile Pro Phe Leu Ile
Arg Gln Lys Ile Ile Ser Ser Leu Asp Pro Pro Cys Arg Arg Gly Ala
             820
                                 825
Asp Trp Arg Thr Leu Ala Gln Lys Leu His Leu Asp Ser His Leu Ser
                             840
Phe Phe Ala Ser Lys Pro Ser Pro Thr Ala Met Ile Leu Asn Leu Trp
                        855
                                             860
Glu Ala Arg His Phe Pro Asn Gly Asn Leu Ser Gln Leu Ala Ala Ala
                    870
Val Ala Gly Leu Gly Gln Pro Asp Ala Gly Leu Phe Thr Val Ser Glu
                885
                                    890
Ala Glu Cys
```

<210> 3 <211> 14536 <212> DNA <213> Homo sapiens

<400> 3

ggagttttcc accatgacta ttgccctgct gggttttgcc atattcttgc tccattgtgc 60 gacctgtgag aagcetetag aagggattet etecteetet gettggeaet teacacaete 120 ccattacaat gccaccatct atgaaaattc ttctcccaag acctatgtgg agagcttcga 180 gaaaatgggc atctacctcg cggagccaca gtgggcagtg aggtaccgga tcatctctgg 240 ggatgtggcc aatgtattta aaactgagga gtatgtggtg ggcaacttct gcttcctaag 300 aataaggaca aagagcagca acacagetet tetgaacaga gaggtgegag acagetacae 360 cctcatcatc caagccacag agaagacctt ggagttggaa gctttgaccc gtgtggtggt 420 ccacatectg gaccagaatg acctgaagee tetettetet ccacettegt acagagteae 480 catctctgag gacatgcccc tgaagagccc catctgcaag gtgactgcca cagatgctga 540 tctaggccag aatgctgagt tctattatgc ctttaacaca aggtcagaga tgtttgccat 600 ccatcccacc agcggtgtgg tcactgtggc tgggaagctt aacgtcacct ggcgaggaaa 660 gcatgagete caggtgetag etgtggaceg catgeggaaa atetetgagg gcaatgggtt 720 tggcagcetg gctgcacttg tggttcatgt ggagcctgcc ctcaggaagc ccccagccat 780 tgcttcggtg gtggtgactc caccagacag caatgatggt accacctatg ccactgtact 840 ggtcgatgca aatagctcag gagctgaagt ggagtcagtg gaagttgttg gtggtgaccc 900 tggaaagcac ttcaaagcca tcaagtctta tgcccggagc aatgagttca gtttggtgtc 960 tgtcaaagac atcaactgga tggagtacct tcatgggttc aacctcagcc tccaggccag 1020 gagtgggagc ggcccttatt tttattccca gatcaggggc tttcacctac caccttccaa 1080 actgtcttcc ctcaaattcg agaaggctgt ttacagagtg cagcttagtg agttttcccc 1140 tectggeage egegtggtga tggtgagagt caceceagee ttececaace tgeagtatgt 1200 tctaaagcca tcttcagaga atgtaggatt taaacttaat gctcgaactg ggttgatcac 1260 caccacaaag ctcatggact tccacgacag agcccactat cagctacaca tcagaacctc 1320 accgggccag gcctccaccg tggtggtcat tgacattgtg gactgcaaca accatgcccc 1380

```
cctcttcaac aggtcttcct atgatggtac cttggatgag aacatccctc caggcaccag 1440
 tgttttggct gtgactgcca ctgaccggga tcatggggaa aatggatatg tcacctattc 1500
 cattgctgga ccaaaagctt tgccattttc tattgacccc tacctgggga tcatctccac 1560
 ctccaaaccc atggactatg aactcatgaa aagaatttat accttccggg taagagcatc 1620
 agactgggga tccccttttc gccgggagaa ggaagtgtcc atttttcttc agctcaggaa 1680
 cttgaatgac aaccagccta tgtttgaaga agtcaactgt acagggtcta tccgccaaga 1740
 ctggccagta gggaaatcga taatgactat gtcagccata gatgtggatg agcttcagaa 1800
 cctaaaatac gagattgtat caggcaatga actagagtat tttgatctaa atcatttctc 1860
 cggagtgata tccctcaaac gcccttttat caatcttact gctggtcaac ccaccagtta 1920
 ttccctgaag attacagcct cagatggcaa aaactatgcc tcacccacaa ctttgaatat 1980
 tactgtggtg aaggaccctc attttgaagt tcctgtaaca tgtgataaaa caggggtatt 2040
 gacacaattc acaaagacta tcctccactt tattgggctt cagaaccagg agtccagtga 2100
 tgaggaattc acttctttaa gcacatatca gattaatcat tacaccccac agtttgagga 2160
 ccacttcccc caatccattg atgtccttga gagtgtccct atcaacaccc ccttggcccg 2220
 cctagcagcc actgaccctg atgctggttt taatggcaaa ctggtctatg tgattgcaga 2280
 tggcaatgag gagggctgct ttgacataga gctggagaca gggctgctca ctgtagctgc 2340
 tcccttggac tatgaagcca ccaatttcta catcctcaat gtaacagtat atgacctggg 2400
 cacaccccag aagtcctcct ggaagctgct gacagtgaat gtgaaagact ggaatgacaa 2460
 cgcacccaga tttcctcccg gtgggtacca gttaaccatc tcggaggaca cagaagttgg 2520
 aaccacaatt gcagagctga caaccaaaga tgctgactcg gaagacaatg gcagggttcg 2580
 ctacaccctg ctaagtccca cagagaagtt ctccctccac cctctcactg gggaactggt 2640
 tgttacagga cacctggacc gcgaatcaga gcctcggtac atactcaagg tggaggccag 2700
 ggatcagece ageaaaggee accagetett etetgteact gaeetgataa teacattgga 2760
ggatgtcaac gacaactctc cccagtgcat cacagaacac aacaggctga aggttccaga 2820
ggacctgccc cccgggactg tcttgacatt tctggatgcc tctgatcctg acctgggccc 2880
cgcaggtgaa gtgcgatatg ttctgatgga tggcgcccat gggaccttcc gggtggacct 2940
gatgacaggg gcgctcattc tggagagaga gctggacttt gagaggcgag ctgggtacaa 3000
tctgagcctg tgggccagtg atggtgggag gcccctagcc cgcaggactc tctgccatgt 3060
ggaggtgatc gtcctggatg tgaatgagaa tctccaccct ccccactttg cctccttcgt 3120
gcaccagggc caggtgcagg agaacageee etegggaact caggtgattg tagtggetgc 3180
ccaggacgat gacagtggct tggatgggga gctccagtac ttcctgcgtg ctggcactgg 3240
actogoageo ttoagoatoa accaagatao aggaatgatt cagactotgg caccootgga 3300
ccgagaattt gcatcttact actggttgac ggtattagca gtggacaggg gttctgtgcc 3360
cctctcttct gtaactgaag tctacatcga ggttacggat gccaatgaca acccacccca 3420
gatgtcccaa gctgtgttct acccctccat ccaggaggat gctcccgtgg gcacctctgt 3480
getteaactg gatgeetggg acceagacte cagetecaaa gggaagetga cetteaacat 3540
caccagtggg aactacatgg gattetttat gatteacect gttacaggte teetatetae 3600
agcccagcag ctggacagag agaacaagga tgaacacatc ctggaggtga ctgtgctgga 3660
caatggggaa ccctcactga agtccacete cagggtggtg gtaggcatet tggacgtcaa 3720
tgacaatcca cctatattct cccacaagct cttcaatgtc cgccttccag agaggctgag 3780
ccctgtgtcc cctgggcctg tgtacaggct ggtggcttca gacctggatg agggtcttaa 3840
tggcagagtc acctacagta tcgaggacag ctatgaggag gccttcagta tcgacctggt 3900
cacaggtgtg gtttcatcca acagcacttt tacagctgga gagtacaaca tcctaacgat 3960
caaggcaaca gacagtgggc agccaccact ctcagccagt gtccggctac acattgagtg 4020
gatecettgg ecceggeegt ectecateee tetggeettt gatgagaeet actaeagett 4080
tacggtcatg gagacggacc ctgtgaacca catggtgggg gtcatcagcg tagagggcag 4140
acceggaete ttetggttea acateteagg tggggataag gacatggaet ttgacattga 4200
gaagaccaca ggcagcatcg tcattgccag gcctcttgat accaggagaa ggtcgaacta 4260
taacttgact gttgaggtga cagatgggtc ccgcaccatt gccacacagg tccacatctt 4320
catgattgcc aacattaacc accatcggcc ccagtttctg gaaactcgtt atgaagtcag 4380
agttececag gacacegtge caggggtaga geteetgega gtecaggeea tagateaaga 4440
caagggcaaa agcctcatct ataccataca tggcagccaa gacccaggaa gtgccagcct 4500
cttccagctg gacccaagca gtggtgtcct ggtaacggtg ggaaaattgg acctcggctc 4560
ggggccctcc cagcacacac tgacagtcat ggtccgagac caggaaatac ctatcaagag 4620
gaacttegtg tgggtgacca tteatgtgga ggatggaaac etecaeceae eeegetteae 4680
tcagctccat tatgaggcaa gtgttcctga caccatagcc cccggcacag agctgctgca 4740
ggtccgagcc atggatgctg accggggagt caatgctgag gtccactact ccctcctgaa 4800
```

```
agggaacagc gaaggtttct tcaacatcaa tgccctgcta ggcatcatta ctctagctca 4860
 aaagettgat caggeaaate atgeeccaea taetetgaea gtgaaggeag aagateaagg 4920
 ctccccacaa tggcatgacc tggctacagt gatcattcat gtctatccct cagataggag 4980
 tgcccccatc ttttcaaaat ctgagtactt tgtagagatc cctgaatcaa tccctgttgg 5040
 ttccccaatc ctccttgtct ctgctatgag cccctctgaa gttacctatg agttaagaga 5100
 gggaaataag gatggagtet tetetatgaa eteatattet ggeettattt eeacceagaa 5160
 gaaattggac catgagaaaa tctcgtctta ccagctgaaa atccgaggca gcaatatggc 5220
 aggtgcattt actgatgtca tggtggtggt tgacataatt gatgaaaatg acaatgctcc 5280
 tatgttctta aagtcaactt ttgtgggcca aattagtgaa gcagctccac tgtatagcat 5340
 gatcatggat aaaaacaaca acccctttgt gattcatgcc tctgacagtg acaaagaagc 5400
 taattccttg ttggtctata aaattttgga gccggaggcc ttgaagtttt tcaaaattga 5460
 tcccagcatg ggaaccctaa ccattgtatc agagatggat tatgagagca tgccctcttt 5520
 ccaattctgt gtctatgtcc atgaccaagg aagccctgta ttatttgcac ccagacctgc 5580
 ccaagtcatc attcatgtca gagatgtgaa tgattcccct cccagattct cagaacagat 5640
 atatgaggta gcaatagtcg ggcctatcca tccaggcatg gagcttctca tggtgcgggc 5700
 cagcgatgaa gactcagaag tcaattatag catcaaaact ggcaatgctg atgaagctgt 5760
 taccatccat cctgtcactg gtagcatatc tgtgctgaat cctgctttcc tgggactctc 5820
 tcggaagete accatcaggg ettetgatgg ettgtateaa gacaetgege tggtaaaaat 5880
ttctttgacc caagtgcttg acaaaagctt gcagtttgat caggatgtct actgggcagc 5940
 tgtgaaggag aacttgcagg acagaaaggc actggtgatt cttggtgccc agggcaatca 6000
tttgaatgac accettteet actttetett gaatggeaca gatatgttte atatggteea 6060
gtcagcaggt gtgttgcaga caagaggtgt ggcgtttgac cgggagcagc aggacactca 6120
tgagttggca gtggaagtga gggacaatcg gacacctcag cgggtggctc agggtttggt 6180
cagagtetet attgaggatg teaatgaeaa tecececaaa tttaageate tgeeetatta 6240
cacaatcatc caagatggca cagagccagg ggatgtcctc tttcaggtat ctgccactga 6300
tgaggacttg gggacaaatg gggctgttac atatgaattt gcagaagatt acacatattt 6360
ccgaattgac ccctatcttg gggacatatc actcaagaaa ccctttgatt atcaagcttt 6420
aaataaatat cacctcaaag tcattgctcg ggatggagga acgccatccc tccagagtga 6480
ggaagaggta cttgtcactg tgagaaataa atccaaccca ctgtttcaga gtccttatta 6540
caaagtcaga gtacctgaaa atatcaccct ctatacccca attctccaca cccaggcccg 6600
gagtecagag ggaeteegge teatetaeaa eattgtggag gaagaaeeet tgatgetgtt 6660
caccactgac ttcaagactg gtgtcctaac agtaacaggg cctttggact atgagtccaa 6720
gaccaaacat gtgttcacag tcagagccac ggatacagct ctggggtcat tttctgaagc 6780
cacagtggaa gtcctagtgg aggatgtcaa tgataaccct cccacttttt cccaattggt 6840
ctataccact tccatctcag aaggettgee tgctcagacc cctgtgatcc aactgttggc 6900
ttctgaccag gactcagggc ggaaccgtga cgtctcttat cagattgtgg aggatggctc 6960
agatgtttcc aagttcttcc agatcaatgg gagcacaggg gagatgtcca cagttcaaga 7020
actggattat gaagcccaac aacactttca tgtgaaagtc agggccatgg ataaaggaga 7080
tcccccactc actggtgaaa cccttgtggt tgtcaatgtg tctgatatca atgacaaccc 7140
cccagagttc agacaacctc aatatgaagc caatgtcagt gaactggcaa cctgtggaca 7200
cctggttctt aaagtccagg ctattgaccc tgacagcaga gacacctccc gcctggagta 7260
cctgattctt tctggcaatc aggacaggca cttcttcatt aacagctcat cgggaataat 7320
ttctatgttc aacctttgca aaaagcacct ggactcttct tacaatttga gggtaggtgc 7380
ttotgatgga gtottoogag caactgtgoo tgtgtacato aacactacaa atgocaacaa 7440
gtacageeca gagttecage ageaeettta tgaggeagaa ttageagaga atgeaatggt 7500
tggaaccaag gtgattgatt tgctagccat agacaaagat agtggtccct atggcactat 7560
agattatact atcatcaata aactagcaag tgagaagttc tccataaacc ccaatggcca 7620
gattgccact ctgcagaaac tggatcggga aaattcaaca gagagagtca ttgctattaa 7680
ggtcatggct cgggatggag gaggaagagt agccttctgc acggtgaaga tcatcctcac 7740
agatgaaaat gacaaccccc cacagttcaa agcatctgag tacacagtat ccattcaatc 7800
caatgtcagt aaagactctc cggttatcca ggtgttggcc tatgatgcag atgaaggtca 7860
gaacgcagat gtcacctact cagtgaaccc agaggaccta gttaaagatg tcattgaaat 7920
taacccagtc actggtgtgg tcaaggtgaa agacagcctg gtgggattgg aaaatcagac 7980
cettgaette tteatcaaag eccaagatgg aggeeeteet eactggaact etetggtgee 8040
agtacgactt caggtggttc ctaaaaaagt atccttaccg aaattttctg aacctttgta 8100
tactttctct gcacctgaag accttccaga ggggtctgaa attgggattg ttaaagcagt 8160
ggcagctcaa gatccagtca tctacagtct agtgcggggc actacacctg agagcaacaa 8220
```

```
ggatggtgtc ttctccctag acccagacac aggggtcata aaggtgagga agcccatgga 8280
 ccacgaatcc accaaattgt accagattga tgtgatggca cattgccttc agaacactga 8340
 tgtggtgtcc ttggtctctg tcaacatcca agtgggagac gtcaatgaca ataggcctgt 8400
 atttgaggct gatccatata aggctgtcct cactgagaat atgccagtgg ggacctcagt 8460
 cattcaagtg actgccattg acaaggacac tgggagagat ggccaggtga gctacaggct 8520
 gtctgcagac cctggtagca atgtccatga gctctttgcc attgacagtg agagtggttg 8580
 gatcaccaca ctccaggaac ttgactgtga gacctgccag acttatcatt ttcatgtggt 8640
 ggcctatgac cacggacaga ccatccagct atcctctcag gccctggttc aggtctccat 8700
 tacagatgag aatgacaatg ctccccgatt tgcttctgaa gagtacagag gatctgtggt 8760
 tgagaacagt gagcctggcg aactggtggc gactctaaag accctggatg ctgacatttc 8820
 tgagcagaac aggcaggtca cctgctacat cacagaggga gaccccctgg gccagtttgg 8880
 catcagccaa gttggagatg agtggaggat ttcctcaagg aagaccctgg accgcgagca 8940
 tacagecaag taettgetea gagteaeage atetgatgge aagtteeagg etteggteae 9000
 tgtggagatc tttgtcctgg acgtcaatga taacagccca cagtgttcac agcttctcta 9060
 tactggcaag gttcatgaag atgtatttcc aggacacttc attttgaagg tttctgccac 9120
agacttggac actgatacca atgctcagat cacatattct ctgcatggcc ctggggcgca 9180
tgaattcaag ctggatcctc atacagggga gctgaccaca ctcactgccc tagaccgaga 9240
aaggaaggat gtgttcaacc ttgttgccaa ggcgacggat ggaggtggcc gatcgtgcca 9300
ggcagacatc accetecatg tggaggatgt gaatgacaat geeegggt tetteeceag 9360
ccactgtgct gtggctgtct tcgacaacac cacagtgaag acccctgtgg ctgtagtatt 9420
tgcccgggat cccgaccaag gcgccaatgc ccaggtggtt tactctctgc cggattcagc 9480
cgaaggccac ttttccatcg acgccaccac gggggtgatc cgcctggaaa agccgctgca 9540
ggtcaggccc caggcaccac tggagctcac ggtccgtgcc tctgacctgg gcaccccaat 9600
accgctgtcc acgctgggca ccgtcacagt ctcggtggtg ggcctagaag actacctgcc 9660
cgtgttcctg aacaccgagc acagcgtgca ggtgcccgag gacgccccac ctggcacgga 9720
ggtgctgcag ctggccaccc tcactcgccc gggcgcagag aagaccggct accgcgtggt 9780
cagegggaac gageaaggea ggtteegeet ggatgetege acagggatee tgtatgteaa 9840
cgcaagcctg gactttgaga caagccccaa gtacttcctg tccattgagt gcagccggaa 9900
gageteetet teeeteagtg aegtgaeeae agteatggte aacateaetg atgteaatga 9960
acaccggccc caattccccc aagatccata tagcacaagg gtcttagaga atgcccttgt 10020
gggtgacgtc atcctcacgg tatcagcgac tgatgaagat ggacccctaa atagtgacat 10080
tacctatagc ctcataggag ggaaccagct tgggcacttc accattcacc ccaaaaaggg 10140
ggagctacag gtggccaagg ccctggaccg ggaacaggcc tctagttatt ccctgaagct 10200
ccgagccaca gacagtgggc agcctccact gcatgaggac acagacatcg ctatccaagt 10260
ggctgatgtc aatgataacc caccgagatt cttccagctc aactacagca ccactgtcca 10320
ggagaactcc cccattggca gcaaagtcct gcagctgatc ctgagtgacc cagattctcc 10380
agagaatggc cccccctact cgtttcgaat caccaagggg aacaacggct ctgccttccg 10440
agtgaccccg gatggatggc tggtgactgc tgagggccta agcaggaggg ctcaggaatg 10500
gtatcagctt cagatccagg cgtcagacag tggcatccct cccctctcgt ctttgacgtc 10560
tgtccgtgtc catgtcacag agcagagcca ctatgcacct tctgctctcc cactggagat 10620
cttcatcact gttggagagg atgagttcca gggtggcatg gtgggtaaga tccatgccac 10680
agaccgagac ccccaggaca cgctgaccta tagcctggca gaagaggaga ccctgggcag 10740
gcacttetea gtgggtgege etgatggeaa gattategee geecagggee tgeetegtgg 10800
ccactactcg ttcaacgtca cggtcagcga tgggaccttc accacgactg ctggggtcca 10860
tgtgtacgtg tggcatgtgg ggcaggaggc tctgcagcag gccatgtgga tgggcttcta 10920
ccagctcacc cccgaggagc tggtgagtga ccactggcgg aacctgcaga ggttcctcag 10980
ccataagctg gacatcaaac gggctaacat tcacttggcc agcctccagc ctgcagaggc 11040
cgtggctggt gtggatgtgc tcctggtctt tgaggggcat tctggaacct tctacgagtt 11100
tcaggagcta gcatccatca tcactcactc agccaaggag atggagcatt cagtggggt 11160
tcagatgcgg tcagctatgc ccatggtgcc ctgccagggg ccaacctgcc agggtcaaat 11220
ctgccataac acagtgcatc tggaccccaa ggttgggccc acgtacagca ccgccaggct 11280
cagcatecta acceegege accaeetgea gaggagetge teetgeaatg gtactgetae 11340
aaggttcagt ggtcagaget atgtgeggta cagggeecca geggetegga actggeacat 11400
ccatttctat ctgaaaacac tccagccaca ggccattctt ctattcacca atgaaacagc 11460
gtccgtctcc ctgaagctgg ccagtggagt gccccagctg gaataccact gtctgggtgg 11520
tttctatgga aacctttcct cccagcgcca tgtgaatgac cacgagtggc actccatcct 11580
ggtggaggag atggacgctt ccattegeet gatggttgac ageatgggea acacetecet 11640
```

```
tgtggtccca gagaactgcc gtggtctgag gcccgaaagg cacctcttgc tgggcggcct 11700
 cattetgttg cattetteet egaatgtete eeagggettt gaaggetgee tggatgetgt 11760
 cgtggtcaac gaagaggctc tagatctgct ggcccctggc aagacggtgg caggcttgct 11820
ggagacacaa gccctcaccc agtgctgcct ccacagtgac tactgcagcc agaacacatg 11880
cctcaatggt gggaagtgct catggaccca tggggcaggc tatgtctgca aatgtccccc 11940
acagttetet gggaageact gtgaacaagg aagggagaac tgtaettttg caccetgeet 12000
ggaaggtgga acttgcatcc tctcccccaa aggagcttcc tgtaactgcc ctcatcctta 12060
cacaggagac aggtgtgaaa tggaggcgag gggttgttca gaaggacact gcctagtcac 12120
tcccgagatc caaagggggg actgggggca gcaggagtta ctgatcatca cagtggccgt 12180
ggcgttcatt atcataagca ctgtcgggct tctcttctac tgccgccgtt gcaagtctca 12240
caageetgtg gecatggagg acceagacet cetggecagg agtgttggtg ttgacaceca 12300
agecatgeet gecategage teaacceatt gagtgecage teetgeaaca aceteaacca 12360
accggaaccc agcaaggcct ctgttccaaa tgaactcgtc acatttggac ccaattctaa 12420
gcaacggcca gtggtctgca gtgtgccccc cagactcccg ccagctgcgg tcccttccca 12480
ctctgacaat gagcctgtca ttaagagaac ctggtccagc gaggagatgg tgtaccctgg 12540
cggagccatg gtctggcccc ctacttactc caggaacgaa cgctgggaat acccccactc 12600
cgaagtgact cagggccctc tgccgccctc ggctcaccgc cactcaaccc cagtcgtgat 12660
gccagagcct aatggcctct atgggggctt ccccttcccc ctggagatgg aaaacaagcg 12720
ggcacctete ecaccegtt acagcaacca gaacctggaa gatetgatge eeteteggee 12780
ccctagtccc cgggagcgcc tggttgcccc ctgtctcaat gagtacacgg ccatcagcta 12840
ctaccactcg cagttccggc agggagggg agggccctgc ctggcagacg ggggctacaa 12900
gggggtgggt atgcgcctca gccgagctgg gccctcttat gctgtctgtg aggtggaggg 12960
ggcacctctt gcaggccagg gccagccccg ggtgcccccc aactatgagg gctctgacat 13020
ggtggagagt gattatggca gctgtgagga ggtcatgttc tagcttccca ttcccagagc 13080
aaggcaggcg ggaggccaag gactggactt ggcttatttc ttcctgtctc gtagggggtg 13140
agttgagtgt ggctgggaga gtgggaggga agccctcagc ccaggctgtt gtcccttgaa 13200
atgtgctctt ccaatccccc acctagtccc tgagggtgga gggaagctga ggatagagct 13260
ccagaaacag cactagggtc ccaggagagg ggcatttcta gagcagtgac cctggaaaac 13320
caggaacaat tgactcctgg ggtgggcgac agacaggagg gctccctgat ctgccggctc 13380
tcagtccccg gggcaaagcc tgattgactg tgctggctca acttcaccaa gatgcattct 13440
catacctgcc cacageteca ttttggagge aggeaggttg gtgeetgaca gacaaccaet 13500
acgegggeeg tacagaggag etetagaggg etgegtggea teeteetagg ggetgagagg 13560
tgagcagcag gggagcgggc acagtcccct ctgcccctgc ctcagtcgag cactcactgt 13620
gtctttgtca agtgtctgct ccacgtcagg cactgtgctt tgcaccgggg agaaaatggt 13680
gatggagggc aacaaggact cegaggagca ceaccaggec tegggeecea gaggteeegc 13740
teeteageet acaegeagag gaaegggeee aceteagagt caeaceaetg getgeeagte 13800
agggcctgcc aggagtctac acagctctga accttctttg ttaaagaatt cagacctcat 13860
ggaactctgg gttcttcatc ccaagtttcc caggcacttt tggccaaagg aaggaaggaa 13920
ctaattcttc attttaaaaa ttcttaggca ctttttgacc ttgctgtctg gatgagtttc 13980
ctcaatggga tttttcttcc ctagacacaa ggaagtctga actcctattt agggccggtt 14040
ggaagcaggg agctggaccg cagtgtccag gctggacacc tgccattgcc tcctctccac 14100
tgcagacgcc tgcccatcaa gtattacctg cagcgactca accctatgca tggagggtca 14160
atgtgggcac atgtctacac atgtgggtgc ccatggatag tacgtgtgta cacatgtgta 14220
gagtgtatgt agccaggagt ggtggggacc agaagcctct gtggcctttg gtgacctcac 14280
cactccctcc cacccagtcc ctccctctgg tccactgcct tttcatatgt gttgtttctg 14340
gagacagaag tcaaaaggaa gagcagtgga gccttgccca cagggctgct gcttcatgcg 14400
agagggagat gtgtgggcga gagccaattt gtgtgagtgg tttgtggctg tgtgtgtgac 14460
tgtgagtgtg agtgacagat acatagtttc attggtcatt tttttttta acaataaagt 14520
atctttttt actgtt
```

```
<210> 4
```

<sup>&</sup>lt;211> 4349

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapiens

<sup>&</sup>lt;400> 4

- Met Thr Ile Ala Leu Leu Gly Phe Ala Ile Phe Leu Leu His Cys Ala 1 5 10 15
- Thr Cys Glu Lys Pro Leu Glu Gly Ile Leu Ser Ser Ser Ala Trp His
  20 25 30
- Phe Thr His Ser His Tyr Asn Ala Thr Ile Tyr Glu Asn Ser Ser Pro 35 40 45
- Lys Thr Tyr Val Glu Ser Phe Glu Lys Met Gly Ile Tyr Leu Ala Glu 50 55 60
- Pro Gln Trp Ala Val Arg Tyr Arg Ile Ile Ser Gly Asp Val Ala Asn
  65 70 75 80
- Val Phe Lys Thr Glu Glu Tyr Val Val Gly Asn Phe Cys Phe Leu Arg 85 90 95
- Ile Arg Thr Lys Ser Ser Asn Thr Ala Leu Leu Asn Arg Glu Val Arg
  100 105 110
- Asp Ser Tyr Thr Leu Ile Ile Gln Ala Thr Glu Lys Thr Leu Glu Leu 115 120 125
- Glu Ala Leu Thr Arg Val Val Val His Ile Leu Asp Gln Asn Asp Leu 130 135 140
- Lys Pro Leu Phe Ser Pro Pro Ser Tyr Arg Val Thr Ile Ser Glu Asp 145 150 155 160
- Met Pro Leu Lys Ser Pro Ile Cys Lys Val Thr Ala Thr Asp Ala Asp 165 170 175
- Leu Gly Gln Asn Ala Glu Phe Tyr Tyr Ala Phe Asn Thr Arg Ser Glu 180 185 190
- Met Phe Ala Ile His Pro Thr Ser Gly Val Val Thr Val Ala Gly Lys 195 200 205
- Leu Asn Val Thr Trp Arg Gly Lys His Glu Leu Gln Val Leu Ala Val 210 215 220
- Asp Arg Met Arg Lys Ile Ser Glu Gly Asn Gly Phe Gly Ser Leu Ala 225 230 235 240
- Ala Leu Val Val His Val Glu Pro Ala Leu Arg Lys Pro Pro Ala Ile 245 250 255
- Ala Ser Val Val Val Thr Pro Pro Asp Ser Asn Asp Gly Thr Thr Tyr 260 265 270
- Ala Thr Val Leu Val Asp Ala Asn Ser Ser Gly Ala Glu Val Glu Ser 275 280 285
- Val Glu Val Val Gly Gly Asp Pro Gly Lys His Phe Lys Ala Ile Lys 290 295 300

Ser Tyr Ala Arg Ser Asn Glu Phe Ser Leu Val Ser Val Lys Asp Ile Asn Trp Met Glu Tyr Leu His Gly Phe Asn Leu Ser Leu Gln Ala Arg 330 Ser Gly Ser Gly Pro Tyr Phe Tyr Ser Gln Ile Arg Gly Phe His Leu Pro Pro Ser Lys Leu Ser Ser Leu Lys Phe Glu Lys Ala Val Tyr Arg 360 Val Gln Leu Ser Glu Phe Ser Pro Pro Gly Ser Arg Val Val Met Val Arg Val Thr Pro Ala Phe Pro Asn Leu Gln Tyr Val Leu Lys Pro Ser 390 395 Ser Glu Asn Val Gly Phe Lys Leu Asn Ala Arg Thr Gly Leu Ile Thr Thr Thr Lys Leu Met Asp Phe His Asp Arg Ala His Tyr Gln Leu His Ile Arg Thr Ser Pro Gly Gln Ala Ser Thr Val Val Val Ile Asp Ile 440 Val Asp Cys Asn Asn His Ala Pro Leu Phe Asn Arg Ser Ser Tyr Asp Gly Thr Leu Asp Glu Asn Ile Pro Pro Gly Thr Ser Val Leu Ala Val 475 Thr Ala Thr Asp Arg Asp His Gly Glu Asn Gly Tyr Val Thr Tyr Ser 485 Ile Ala Gly Pro Lys Ala Leu Pro Phe Ser Ile Asp Pro Tyr Leu Gly 505 Ile Ile Ser Thr Ser Lys Pro Met Asp Tyr Glu Leu Met Lys Arg Ile 520 Tyr Thr Phe Arg Val Arg Ala Ser Asp Trp Gly Ser Pro Phe Arg Arg Glu Lys Glu Val Ser Ile Phe Leu Gln Leu Arg Asn Leu Asn Asp Asn 550 555 Gln Pro Met Phe Glu Glu Val Asn Cys Thr Gly Ser Ile Arg Gln Asp 565 Trp Pro Val Gly Lys Ser Ile Met Thr Met Ser Ala Ile Asp Val Asp 585 Glu Leu Gln Asn Leu Lys Tyr Glu Ile Val Ser Gly Asn Glu Leu Glu 600 605

Tyr Phe Asp Leu Asn His Phe Ser Gly Val Ile Ser Leu Lys Arg Pro 610 Phe Ile Asn Leu Thr Ala Gly Gln Pro Thr Ser Tyr Ser Leu Lys Ile 630 Thr Ala Ser Asp Gly Lys Asn Tyr Ala Ser Pro Thr Thr Leu Asn Ile 645 650 Thr Val Val Lys Asp Pro His Phe Glu Val Pro Val Thr Cys Asp Lys 665 Thr Gly Val Leu Thr Gln Phe Thr Lys Thr Ile Leu His Phe Ile Gly 680 Leu Gln Asn Gln Glu Ser Ser Asp Glu Glu Phe Thr Ser Leu Ser Thr Tyr Gln Ile Asn His Tyr Thr Pro Gln Phe Glu Asp His Phe Pro Gln 710 Ser Ile Asp Val Leu Glu Ser Val Pro Ile Asn Thr Pro Leu Ala Arg 725 730 Leu Ala Ala Thr Asp Pro Asp Ala Gly Phe Asn Gly Lys Leu Val Tyr 745 Val Ile Ala Asp Gly Asn Glu Glu Gly Cys Phe Asp Ile Glu Leu Glu Thr Gly Leu Leu Thr Val Ala Ala Pro Leu Asp Tyr Glu Ala Thr Asn 775 Phe Tyr Ile Leu Asn Val Thr Val Tyr Asp Leu Gly Thr Pro Gln Lys 790 795 Ser Ser Trp Lys Leu Leu Thr Val Asn Val Lys Asp Trp Asn Asp Asn 810 Ala Pro Arg Phe Pro Pro Gly Gly Tyr Gln Leu Thr Ile Ser Glu Asp 820 Thr Glu Val Gly Thr Thr Ile Ala Glu Leu Thr Thr Lys Asp Ala Asp 840 Ser Glu Asp Asn Gly Arg Val Arg Tyr Thr Leu Leu Ser Pro Thr Glu 855 Lys Phe Ser Leu His Pro Leu Thr Gly Glu Leu Val Val Thr Gly His 865 870 875 Leu Asp Arg Glu Ser Glu Pro Arg Tyr Ile Leu Lys Val Glu Ala Arg 890 Asp Gln Pro Ser Lys Gly His Gln Leu Phe Ser Val Thr Asp Leu Ile 900 905 910

- Ile Thr Leu Glu Asp Val Asn Asp Asn Ser Pro Gln Cys Ile Thr Glu 915 920 925
- His Asn Arg Leu Lys Val Pro Glu Asp Leu Pro Pro Gly Thr Val Leu 930 935 940
- Thr Phe Leu Asp Ala Ser Asp Pro Asp Leu Gly Pro Ala Gly Glu Val 945 950 955 960
- Arg Tyr Val Leu Met Asp Gly Ala His Gly Thr Phe Arg Val Asp Leu 965 970 975
- Met Thr Gly Ala Leu Ile Leu Glu Arg Glu Leu Asp Phe Glu Arg Arg 980 985 990
- Ala Gly Tyr Asn Leu Ser Leu Trp Ala Ser Asp Gly Gly Arg Pro Leu 995 1000 1005
- Ala Arg Arg Thr Leu Cys His Val Glu Val Ile Val Leu Asp Val Asn 1010 1015 1020
- Glu Asn Leu His Pro Pro His Phe Ala Ser Phe Val His Gln Gly Gln 1025 1030 1035 1040
- Val Gln Glu Asn Ser Pro Ser Gly Thr Gln Val Ile Val Val Ala Ala 1045 1050 1055
- Gln Asp Asp Ser Gly Leu Asp Gly Glu Leu Gln Tyr Phe Leu Arg 1060 1065 1070
- Ala Gly Thr Gly Leu Ala Ala Phe Ser Ile Asn Gln Asp Thr Gly Met 1075 1080 1085
- Ile Gln Thr Leu Ala Pro Leu Asp Arg Glu Phe Ala Ser Tyr Tyr Trp 1090 1095 1100
- Leu Thr Val Leu Ala Val Asp Arg Gly Ser Val Pro Leu Ser Ser Val 1105 1110 1115 1120
- Thr Glu Val Tyr Ile Glu Val Thr Asp Ala Asn Asp Asn Pro Pro Gln
  1125 1130 1135
- Met Ser Gln Ala Val Phe Tyr Pro Ser Ile Gln Glu Asp Ala Pro Val 1140 1145 1150
- Gly Thr Ser Val Leu Gln Leu Asp Ala Trp Asp Pro Asp Ser Ser 1155 1160 1165
- Lys Gly Lys Leu Thr Phe Asn Ile Thr Ser Gly Asn Tyr Met Gly Phe 1170 1180
- Phe Met Ile His Pro Val Thr Gly Leu Leu Ser Thr Ala Gln Gln Leu 1185 1190 1195 1200
- Asp Arg Glu Asn Lys Asp Glu His Ile Leu Glu Val Thr Val Leu Asp 1205 1210 1215

- Asn Gly Glu Pro Ser Leu Lys Ser Thr Ser Arg Val Val Gly Ile 1220 1225 1230
- Leu Asp Val Asn Asp Asn Pro Pro Ile Phe Ser His Lys Leu Phe Asn 1235 1240 1245
- Val Arg Leu Pro Glu Arg Leu Ser Pro Val Ser Pro Gly Pro Val Tyr 1250 1255 1260
- Arg Leu Val Ala Ser Asp Leu Asp Glu Gly Leu Asn Gly Arg Val Thr 1265 1270 1275 1280
- Tyr Ser Ile Glu Asp Ser Tyr Glu Glu Ala Phe Ser Ile Asp Leu Val 1285 1290 1295
- Thr Gly Val Val Ser Ser Asn Ser Thr Phe Thr Ala Gly Glu Tyr Asn 1300 1305 1310
- Ile Leu Thr Ile Lys Ala Thr Asp Ser Gly Gln Pro Pro Leu Ser Ala 1315 1320 1325
- Ser Val Arg Leu His Ile Glu Trp Ile Pro Trp Pro Arg Pro Ser Ser 1330 1335 1340
- Ile Pro Leu Ala Phe Asp Glu Thr Tyr Tyr Ser Phe Thr Val Met Glu 1345 1350 1355 1360
- Thr Asp Pro Val Asn His Met Val Gly Val Ile Ser Val Glu Gly Arg 1365 1370 1375
- Pro Gly Leu Phe Trp Phe Asn Ile Ser Gly Gly Asp Lys Asp Met Asp 1380 1385 1390
- Phe Asp Ile Glu Lys Thr Thr Gly Ser Ile Val Ile Ala Arg Pro Leu 1395 1400 1405
- Asp Thr Arg Arg Ser Asn Tyr Asn Leu Thr Val Glu Val Thr Asp 1410 1415 1420
- Gly Ser Arg Thr Ile Ala Thr Gln Val His Ile Phe Met Ile Ala Asn 1425 1430 1435 1440
- Ile Asn His His Arg Pro Gln Phe Leu Glu Thr Arg Tyr Glu Val Arg 1445 1450 1455
- Val Pro Gln Asp Thr Val Pro Gly Val Glu Leu Leu Arg Val Gln Ala 1460 1465 1470
- Ile Asp Gln Asp Lys Gly Lys Ser Leu Ile Tyr Thr Ile His Gly Ser 1475 1480 1485
- Gln Asp Pro Gly Ser Ala Ser Leu Phe Gln Leu Asp Pro Ser Ser Gly 1490 1495 1500
- Val Leu Val Thr Val Gly Lys Leu Asp Leu Gly Ser Gly Pro Ser Gln 1505 1510 1515 1520

- His Thr Leu Thr Val Met Val Arg Asp Gln Glu Ile Pro Ile Lys Arg 1525 1530 1535
- Asn Phe Val Trp Val Thr Ile His Val Glu Asp Gly Asn Leu His Pro
  1540 1545 1550
- Pro Arg Phe Thr Gln Leu His Tyr Glu Ala Ser Val Pro Asp Thr Ile 1555 1560 1565
- Ala Pro Gly Thr Glu Leu Leu Gln Val Arg Ala Met Asp Ala Asp Arg 1570 1580
- Gly Val Asn Ala Glu Val His Tyr Ser Leu Leu Lys Gly Asn Ser Glu 1585 1590 1595 1600
- Gly Phe Phe Asn Ile Asn Ala Leu Leu Gly Ile Ile Thr Leu Ala Gln 1605 1610 1615
- Lys Leu Asp Gln Ala Asn His Ala Pro His Thr Leu Thr Val Lys Ala 1620 1625 1630
- Glu Asp Gln Gly Ser Pro Gln Trp His Asp Leu Ala Thr Val Ile Ile 1635 1640 1645
- His Val Tyr Pro Ser Asp Arg Ser Ala Pro Ile Phe Ser Lys Ser Glu 1650 1660
- Tyr Phe Val Glu Ile Pro Glu Ser Ile Pro Val Gly Ser Pro Ile Leu 1665 1670 1680
- Leu Val Ser Ala Met Ser Pro Ser Glu Val Thr Tyr Glu Leu Arg Glu 1685 1690 1695
- Gly Asn Lys Asp Gly Val Phe Ser Met Asn Ser Tyr Ser Gly Leu Ile 1700 1705 1710
- Ser Thr Gln Lys Lys Leu Asp His Glu Lys Ile Ser Ser Tyr Gln Leu 1715 1720 1725
- Lys Ile Arg Gly Ser Asn Met Ala Gly Ala Phe Thr Asp Val Met Val 1730 1735 1740
- Val Val Asp Ile Ile Asp Glu Asn Asp Asn Ala Pro Met Phe Leu Lys 1745 1750 1755 1760
- Ser Thr Phe Val Gly Gln Ile Ser Glu Ala Ala Pro Leu Tyr Ser Met 1765 1770 1775
- Ile Met Asp Lys Asn Asn Pro Phe Val Ile His Ala Ser Asp Ser 1780 1785 1790
- Asp Lys Glu Ala Asn Ser Leu Leu Val Tyr Lys Ile Leu Glu Pro Glu 1795 1800 1805
- Ala Leu Lys Phe Phe Lys Ile Asp Pro Ser Met Gly Thr Leu Thr Ile 1810 1815 1820

- Val Ser Glu Met Asp Tyr Glu Ser Met Pro Ser Phe Gln Phe Cys Val 1825 1830 1835 1840
- Tyr Val His Asp Gln Gly Ser Pro Val Leu Phe Ala Pro Arg Pro Ala 1845 1850 1855
- Gln Val Ile Ile His Val Arg Asp Val Asn Asp Ser Pro Pro Arg Phe 1860 1865 1870
- Ser Glu Gln Ile Tyr Glu Val Ala Ile Val Gly Pro Ile His Pro Gly 1875 1880 1885
- Met Glu Leu Leu Met Val Arg Ala Ser Asp Glu Asp Ser Glu Val Asn 1890 1895 1900
- Tyr Ser Ile Lys Thr Gly Asn Ala Asp Glu Ala Val Thr Ile His Pro 1905 1910 1915 1920
- Val Thr Gly Ser Ile Ser Val Leu Asn Pro Ala Phe Leu Gly Leu Ser 1925 1930 1935
- Arg Lys Leu Thr Ile Arg Ala Ser Asp Gly Leu Tyr Gln Asp Thr Ala 1940 1945 1950
- Leu Val Lys Ile Ser Leu Thr Gln Val Leu Asp Lys Ser Leu Gln Phe 1955 1960 1965
- Asp Gln Asp Val Tyr Trp Ala Ala Val Lys Glu Asn Leu Gln Asp Arg 1970 1975 1980
- Lys Ala Leu Val Ile Leu Gly Ala Gln Gly Asn His Leu Asn Asp Thr 1985 1990 1995 2000
- Leu Ser Tyr Phe Leu Leu Asn Gly Thr Asp Met Phe His Met Val Gln 2005 2010 2015
- Ser Ala Gly Val Leu Gln Thr Arg Gly Val Ala Phe Asp Arg Glu Gln 2020 2025 2030
- Gln Asp Thr His Glu Leu Ala Val Glu Val Arg Asp Asn Arg Thr Pro 2035 2040 2045
- Gln Arg Val Ala Gln Gly Leu Val Arg Val Ser Ile Glu Asp Val Asn 2050 2055 2060
- Asp Asn Pro Pro Lys Phe Lys His Leu Pro Tyr Tyr Thr Ile Ile Gln 2065 2070 2075 2080
- Asp Gly Thr Glu Pro Gly Asp Val Leu Phe Gln Val Ser Ala Thr Asp 2085 2090 2095
- Glu Asp Leu Gly Thr Asn Gly Ala Val Thr Tyr Glu Phe Ala Glu Asp 2100 2105 2110
- Tyr Thr Tyr Phe Arg Ile Asp Pro Tyr Leu Gly Asp Ile Ser Leu Lys 2115 2120 2125

- Lys Pro Phe Asp Tyr Gln Ala Leu Asn Lys Tyr His Leu Lys Val Ile 2130 2135 2140
- Ala Arg Asp Gly Gly Thr Pro Ser Leu Gln Ser Glu Glu Glu Val Leu 2145 2150 2155 2160
- Val Thr Val Arg Asn Lys Ser Asn Pro Leu Phe Gln Ser Pro Tyr Tyr 2165 2170 2175
- Lys Val Arg Val Pro Glu Asn Ile Thr Leu Tyr Thr Pro Ile Leu His 2180 2185 2190
- Thr Gln Ala Arg Ser Pro Glu Gly Leu Arg Leu Ile Tyr Asn Ile Val 2195 2200 2205
- Glu Glu Glu Pro Leu Met Leu Phe Thr Thr Asp Phe Lys Thr Gly Val 2210 2215 2220
- Leu Thr Val Thr Gly Pro Leu Asp Tyr Glu Ser Lys Thr Lys His Val 2225 2230 2235 2240
- Phe Thr Val Arg Ala Thr Asp Thr Ala Leu Gly Ser Phe Ser Glu Ala 2245 2250 2255
- Thr Val Glu Val Leu Val Glu Asp Val Asn Asp Asn Pro Pro Thr Phe 2260 2265 2270
- Ser Gln Leu Val Tyr Thr Thr Ser Ile Ser Glu Gly Leu Pro Ala Gln 2275 2280 2285
- Thr Pro Val Ile Gln Leu Leu Ala Ser Asp Gln Asp Ser Gly Arg Asn 2290 2295 2300
- Arg Asp Val Ser Tyr Gln Ile Val Glu Asp Gly Ser Asp Val Ser Lys 2305 2310 2315 2320
- Phe Phe Gln Ile Asn Gly Ser Thr Gly Glu Met Ser Thr Val Gln Glu 2325 2330 2335
- Leu Asp Tyr Glu Ala Gln Gln His Phe His Val Lys Val Arg Ala Met 2340 2345 2350
- Asp Lys Gly Asp Pro Pro Leu Thr Gly Glu Thr Leu Val Val Val Asn 2355 2360 2365
- Val Ser Asp Ile Asn Asp Asn Pro Pro Glu Phe Arg Gln Pro Gln Tyr 2370 2375 2380
- Glu Ala Asn Val Ser Glu Leu Ala Thr Cys Gly His Leu Val Leu Lys 2385 2390 2395 2400
- Val Gln Ala Ile Asp Pro Asp Ser Arg Asp Thr Ser Arg Leu Glu Tyr 2405 2410 2415
- Leu Ile Leu Ser Gly Asn Gln Asp Arg His Phe Phe Ile Asn Ser Ser 2420 2425 2430

- Ser Gly Ile Ile Ser Met Phe Asn Leu Cys Lys Lys His Leu Asp Ser 2435 2440 2445
- Ser Tyr Asn Leu Arg Val Gly Ala Ser Asp Gly Val Phe Arg Ala Thr 2450 2455 2460
- Val Pro Val Tyr Ile Asn Thr Thr Asn Ala Asn Lys Tyr Ser Pro Glu 2465 2470 2475 2480
- Phe Gln Gln His Leu Tyr Glu Ala Glu Leu Ala Glu Asn Ala Met Val\$2485\$ 2490 2495
- Gly Thr Lys Val Ile Asp Leu Leu Ala Ile Asp Lys Asp Ser Gly Pro 2500 2505 2510
- Tyr Gly Thr Ile Asp Tyr Thr Ile Ile Asn Lys Leu Ala Ser Glu Lys 2515 2520 2525
- Phe Ser Ile Asn Pro Asn Gly Gln Ile Ala Thr Leu Gln Lys Leu Asp 2530 2540
- Arg Glu Asn Ser Thr Glu Arg Val Ile Ala Ile Lys Val Met Ala Arg 2545 2550 2555 2560
- Asp Gly Gly Arg Val Ala Phe Cys Thr Val Lys Ile Ile Leu Thr 2565 2570 2575
- Asp Glu Asn Asp Asn Pro Pro Gln Phe Lys Ala Ser Glu Tyr Thr Val 2580 2585 2590
- Ser Ile Gln Ser Asn Val Ser Lys Asp Ser Pro Val Ile Gln Val Leu 2595 2600 2605
- Ala Tyr Asp Ala Asp Glu Gly Gln Asn Ala Asp Val Thr Tyr Ser Val 2610 2615 2620
- Asn Pro Glu Asp Leu Val Lys Asp Val Ile Glu Ile Asn Pro Val Thr 2625 2630 2635 2640
- Gly Val Val Lys Val Lys Asp Ser Leu Val Gly Leu Glu Asn Gln Thr 2645 2650 2655
- Leu Asp Phe Phe Ile Lys Ala Gln Asp Gly Gly Pro Pro His Trp Asn 2660 2665 2670
- Ser Leu Val Pro Val Arg Leu Gln Val Val Pro Lys Lys Val Ser Leu 2675 2680 2685
- Pro Lys Phe Ser Glu Pro Leu Tyr Thr Phe Ser Ala Pro Glu Asp Leu 2690 2695 2700
- Pro Glu Gly Ser Glu Ile Gly Ile Val Lys Ala Val Ala Ala Gln Asp 2705 2710 2715 2720
- Pro Val Ile Tyr Ser Leu Val Arg Gly Thr Thr Pro Glu Ser Asn Lys 2725 2730 2735

- Asp Gly Val Phe Ser Leu Asp Pro Asp Thr Gly Val Ile Lys Val Arg 2740 2745 2750
- Lys Pro Met Asp His Glu Ser Thr Lys Leu Tyr Gln Ile Asp Val Met 2755 2760 2765
- Ala His Cys Leu Gln Asn Thr Asp Val Val Ser Leu Val Ser Val Asn 2770 2775 2780
- Ile Gln Val Gly Asp Val Asn Asp Asn Arg Pro Val Phe Glu Ala Asp 2785 2790 2795 2800
- Pro Tyr Lys Ala Val Leu Thr Glu Asn Met Pro Val Gly Thr Ser Val 2805 2810 2815
- Ile Gln Val Thr Ala Ile Asp Lys Asp Thr Gly Arg Asp Gly Gln Val 2820 2825 2830
- Ser Tyr Arg Leu Ser Ala Asp Pro Gly Ser Asn Val His Glu Leu Phe 2835 2840 2845
- Ala Ile Asp Ser Glu Ser Gly Trp Ile Thr Thr Leu Gln Glu Leu Asp 2850 2855 2860
- Cys Glu Thr Cys Gln Thr Tyr His Phe His Val Val Ala Tyr Asp His 2865 2870 2875 2880
- Gly Gln Thr Ile Gln Leu Ser Ser Gln Ala Leu Val Gln Val Ser Ile 2885 2890 2895
- Thr Asp Glu Asn Asp Asn Ala Pro Arg Phe Ala Ser Glu Glu Tyr Arg
  2900 2905 2910
- Gly Ser Val Val Glu Asn Ser Glu Pro Gly Glu Leu Val Ala Thr Leu 2915 2920 2925
- Lys Thr Leu Asp Ala Asp Ile Ser Glu Gln Asn Arg Gln Val Thr Cys 2930 2935 2940
- Tyr Ile Thr Glu Gly Asp Pro Leu Gly Gln Phe Gly Ile Ser Gln Val 2945 2950 2955 2960
- Gly Asp Glu Trp Arg Ile Ser Ser Arg Lys Thr Leu Asp Arg Glu His 2965 2970 2975
- Thr Ala Lys Tyr Leu Leu Arg Val Thr Ala Ser Asp Gly Lys Phe Gln 2980 2985 2990
- Ala Ser Val Thr Val Glu Ile Phe Val Leu Asp Val Asn Asp Asn Ser 2995 3000 3005
- Pro Gln Cys Ser Gln Leu Leu Tyr Thr Gly Lys Val His Glu Asp Val 3010 3015 3020
- Phe Pro Gly His Phe Ile Leu Lys Val Ser Ala Thr Asp Leu Asp Thr 3025 3030 3035 3040

- Asp Thr Asn Ala Gln Ile Thr Tyr Ser Leu His Gly Pro Gly Ala His 3045 3050 3055
- Glu Phe Lys Leu Asp Pro His Thr Gly Glu Leu Thr Thr Leu Thr Ala 3060 3065 3070
- Leu Asp Arg Glu Arg Lys Asp Val Phe Asn Leu Val Ala Lys Ala Thr 3075 3080 3085
- Asp Gly Gly Gly Arg Ser Cys Gln Ala Asp Ile Thr Leu His Val Glu 3090 3095 3100
- Asp Val Asn Asp Asn Ala Pro Arg Phe Phe Pro Ser His Cys Ala Val 3105 3110 3115 3120
- Ala Val Phe Asp Asn Thr Thr Val Lys Thr Pro Val Ala Val Val Phe
  3125 3130 3135
- Ala Arg Asp Pro Asp Gln Gly Ala Asn Ala Gln Val Val Tyr Ser Leu 3140 3145 3150
- Pro Asp Ser Ala Glu Gly His Phe Ser Ile Asp Ala Thr Thr Gly Val 3155 3160 3165
- Ile Arg Leu Glu Lys Pro Leu Gln Val Arg Pro Gln Ala Pro Leu Glu 3170 3180
- Leu Thr Val Arg Ala Ser Asp Leu Gly Thr Pro Ile Pro Leu Ser Thr 3185 3190 3195 3200
- Leu Gly Thr Val Thr Val Ser Val Val Gly Leu Glu Asp Tyr Leu Pro 3205 3210 3215
- Val Phe Leu Asn Thr Glu His Ser Val Gln Val Pro Glu Asp Ala Pro 3220 3225 3230
- Pro Gly Thr Glu Val Leu Gln Leu Ala Thr Leu Thr Arg Pro Gly Ala 3235 3240 3245
- Glu Lys Thr Gly Tyr Arg Val Val Ser Gly Asn Glu Gln Gly Arg Phe 3250 3260
- Arg Leu Asp Ala Arg Thr Gly Ile Leu Tyr Val Asn Ala Ser Leu Asp 3265 3270 3280
- Phe Glu Thr Ser Pro Lys Tyr Phe Leu Ser Ile Glu Cys Ser Arg Lys 3285 3290 3295
- Ser Ser Ser Ser Leu Ser Asp Val Thr Thr Val Met Val Asn Ile Thr 3300 3305 3310
- Asp Val Asn Glu His Arg Pro Gln Phe Pro Gln Asp Pro Tyr Ser Thr 3315 3320 3325
- Arg Val Leu Glu Asn Ala Leu Val Gly Asp Val Ile Leu Thr Val Ser 3330 3340

- Ala Thr Asp Glu Asp Gly Pro Leu Asn Ser Asp Ile Thr Tyr Ser Leu 3345 3350 3355 3360
- Ile Gly Gly Asn Gln Leu Gly His Phe Thr Ile His Pro Lys Lys Gly 3365 3370 3375
- Glu Leu Gln Val Ala Lys Ala Leu Asp Arg Glu Gln Ala Ser Ser Tyr 3380 3385 3390
- Ser Leu Lys Leu Arg Ala Thr Asp Ser Gly Gln Pro Pro Leu His Glu 3395 3400 3405
- Asp Thr Asp Ile Ala Ile Gln Val Ala Asp Val Asn Asp Asn Pro Pro 3410 3415 3420
- Arg Phe Phe Gln Leu Asn Tyr Ser Thr Thr Val Gln Glu Asn Ser Pro 3425 3430 3435 3440
- Ile Gly Ser Lys Val Leu Gln Leu Ile Leu Ser Asp Pro Asp Ser Pro 3445 3450 3455
- Glu Asn Gly Pro Pro Tyr Ser Phe Arg Ile Thr Lys Gly Asn Asn Gly 3460 3465 3470
- Ser Ala Phe Arg Val Thr Pro Asp Gly Trp Leu Val Thr Ala Glu Gly 3475 3480 3485
- Leu Ser Arg Arg Ala Gln Glu Trp Tyr Gln Leu Gln Ile Gln Ala Ser 3490 3495 3500
- Asp Ser Gly Ile Pro Pro Leu Ser Ser Leu Thr Ser Val Arg Val His 3505 3510 3515 3520
- Val Thr Glu Gln Ser His Tyr Ala Pro Ser Ala Leu Pro Leu Glu Ile 3525 3530 3535
- Phe Ile Thr Val Gly Glu Asp Glu Phe Gln Gly Gly Met Val Gly Lys 3540 3545 3550
- Ile His Ala Thr Asp Arg Asp Pro Gln Asp Thr Leu Thr Tyr Ser Leu 3555 3560 3565
- Ala Glu Glu Glu Thr Leu Gly Arg His Phe Ser Val Gly Ala Pro Asp 3570 3580
- Gly Lys Ile Ile Ala Ala Gln Gly Leu Pro Arg Gly His Tyr Ser Phe 3585 3590 3595 3600
- Asn Val Thr Val Ser Asp Gly Thr Phe Thr Thr Ala Gly Val His 3605 3610 3615
- Val Tyr Val Trp His Val Gly Gln Glu Ala Leu Gln Gln Ala Met Trp 3620 3625 3630
- Met Gly Phe Tyr Gln Leu Thr Pro Glu Glu Leu Val Ser Asp His Trp 3635 3640 3645

- Arg Asn Leu Gln Arg Phe Leu Ser His Lys Leu Asp Ile Lys Arg Ala 3650 3660
- Asn Ile His Leu Ala Ser Leu Gln Pro Ala Glu Ala Val Ala Gly Val 3665 3670 3675 3680
- Asp Val Leu Leu Val Phe Glu Gly His Ser Gly Thr Phe Tyr Glu Phe 3685 3690 3695
- Gln Glu Leu Ala Ser Ile Ile Thr His Ser Ala Lys Glu Met Glu His 3700 3705 3710
- Ser Val Gly Val Gln Met Arg Ser Ala Met Pro Met Val Pro Cys Gln 3715 3720 3725
- Gly Pro Thr Cys Gln Gly Gln Ile Cys His Asn Thr Val His Leu Asp 3730 3735 3740
- Pro Lys Val Gly Pro Thr Tyr Ser Thr Ala Arg Leu Ser Ile Leu Thr 3745 3750 3755 3760
- Pro Arg His His Leu Gln Arg Ser Cys Ser Cys Asn Gly Thr Ala Thr 3765 3770 3775
- Arg Phe Ser Gly Gln Ser Tyr Val Arg Tyr Arg Ala Pro Ala Ala Arg 3780 3785 3790
- Asn Trp His Ile His Phe Tyr Leu Lys Thr Leu Gln Pro Gln Ala Ile 3795 3800 3805
- Leu Leu Phe Thr Asn Glu Thr Ala Ser Val Ser Leu Lys Leu Ala Ser 3810 3815 3820
- Gly Val Pro Gln Leu Glu Tyr His Cys Leu Gly Gly Phe Tyr Gly Asn 3825 3830 3835 3840
- Leu Ser Ser Gln Arg His Val Asn Asp His Glu Trp His Ser Ile Leu 3845 3850 3855
- Val Glu Glu Met Asp Ala Ser Ile Arg Leu Met Val Asp Ser Met Gly 3860 3865 3870
- Asn Thr Ser Leu Val Val Pro Glu Asn Cys Arg Gly Leu Arg Pro Glu 3875 3880 3885
- Arg His Leu Leu Gly Gly Leu Ile Leu Leu His Ser Ser Ser Asn 3890 3895 3900
- Val Ser Gln Gly Phe Glu Gly Cys Leu Asp Ala Val Val Asn Glu 3905 3910 3915 3920
- Glu Ala Leu Asp Leu Leu Ala Pro Gly Lys Thr Val Ala Gly Leu Leu 3925 3930 3935
- Glu Thr Gln Ala Leu Thr Gln Cys Cys Leu His Ser Asp Tyr Cys Ser 3940 3945 3950

- Gln Asn Thr Cys Leu Asn Gly Gly Lys Cys Ser Trp Thr His Gly Ala 3955 3960 3965
- Gly Tyr Val Cys Lys Cys Pro Pro Gln Phe Ser Gly Lys His Cys Glu 3970 3975 3980
- Gln Gly Arg Glu Asn Cys Thr Phe Ala Pro Cys Leu Glu Gly Gly Thr 3985 3990 3995 4000
- Cys Ile Leu Ser Pro Lys Gly Ala Ser Cys Asn Cys Pro His Pro Tyr 4005 4010 4015
- Thr Gly Asp Arg Cys Glu Met Glu Ala Arg Gly Cys Ser Glu Gly His 4020 4025 4030
- Cys Leu Val Thr Pro Glu Ile Gln Arg Gly Asp Trp Gly Gln Glu
  4035 4040 4045
- Leu Leu Ile Ile Thr Val Ala Val Ala Phe Ile Ile Ile Ser Thr Val 4050 4055 4060
- Gly Leu Leu Phe Tyr Cys Arg Arg Cys Lys Ser His Lys Pro Val Ala 4065 4070 4075 4080
- Met Glu Asp Pro Asp Leu Leu Ala Arg Ser Val Gly Val Asp Thr Gln 4085 4090 4095
- Ala Met Pro Ala Ile Glu Leu Asn Pro Leu Ser Ala Ser Ser Cys Asn 4100 4105 4110
- Asn Leu Asn Gln Pro Glu Pro Ser Lys Ala Ser Val Pro Asn Glu Leu 4115 4120 4125
- Val Thr Phe Gly Pro Asn Ser Lys Gln Arg Pro Val Val Cys Ser Val 4130 4135 4140
- Pro Pro Arg Leu Pro Pro Ala Ala Val Pro Ser His Ser Asp Asn Glu 4145 4150 4155 4160
- Pro Val Ile Lys Arg Thr Trp Ser Ser Glu Glu Met Val Tyr Pro Gly
  4165 4170 4175
- Gly Ala Met Val Trp Pro Pro Thr Tyr Ser Arg Asn Glu Arg Trp Glu 4180 4185 4190
- Tyr Pro His Ser Glu Val Thr Gln Gly Pro Leu Pro Pro Ser Ala His 4195 4200 4205
- Arg His Ser Thr Pro Val Val Met Pro Glu Pro Asn Gly Leu Tyr Gly
  4210 4215 4220
- Gly Phe Pro Phe Pro Leu Glu Met Glu Asn Lys Arg Ala Pro Leu Pro 4225 4230 4235 4240
- Pro Arg Tyr Ser Asn Gln Asn Leu Glu Asp Leu Met Pro Ser Arg Pro
  4245 4250 4255

Pro Ser Pro Arg Glu Arg Leu Val Ala Pro Cys Leu Asn Glu Tyr Thr 4260 4265 4270

Ala Ile Ser Tyr Tyr His Ser Gln Phe Arg Gln Gly Gly Gly Pro 4275 4280 4285

Cys Leu Ala Asp Gly Gly Tyr Lys Gly Val Gly Met Arg Leu Ser Arg 4290 4295 4300

Ala Gly Pro Ser Tyr Ala Val Cys Glu Val Glu Gly Ala Pro Leu Ala 4305 4310 4315 4320

Gly Gln Gly Gln Pro Arg Val Pro Pro Asn Tyr Glu Gly Ser Asp Met 4325 4330 4335

Val Glu Ser Asp Tyr Gly Ser Cys Glu Glu Val Met Phe 4340 4345

<210> 5

<211> 3381

<212> DNA

<213> Homo sapiens

<400> 5

ctagaattca gcggccgctt aattcagaac ggccccctgc cactgccagg aggacggcat 60 catgctgtct gccgactgct ctgagctcgg gctgtccgcc gttccggggg acccggaccc 120 cctgacggct tacctggacc tcagcatgaa caacctcaca gagcttcagc ctggcctctt 180 ccaccacctg cgcttcttgg aggagetgcg tctctctggg aaccatctct cacacatccc 240 aggacaagca ttctctggtc tctacagcct gaaaatcctg atgctqcaga acaatcaqct 300 gggaggaatc cccgcagagg cgctqtqqqa qctqccqaqc ctqcaqtcqc tqcqcctaqa 360 tgccaacctc atctccctgg tcccggagag gagctttgag gggctgtcct ccctccgcca 420 cetetggetg gacgacaatg cacteaegga gatecetgte agggeeetea acaaceteee 480 tgccctgcag gccatgaccc tggccctcaa ccgcatcagc cacatccccg actacgcgtt 540 ccagaatctc accagccttg tggtgctgca tttgcataac aaccgcatcc agcatctggg 600 gacccacage ttegagggge tgeacaatet ggagacaeta gacetgaatt ataacaaget 660 gcaggagttc cctgtggcca tccggaccct gggcagactg caggaactgg ggttccataa 720 caacaacatc aaggccatcc cagaaaaggc cttcatgggg aaccctctgc tacagacgat 780 acacttttat gataacccaa tccagtttgt gggaagatcg qcattccagt acctgcctaa 840 actccacaca ctatctctga atggtgccat ggacatccag gagtttccag atctcaaagg 900 caccaccage etggagatee tgaccetgae eegegeagge ateeggetge teccateggg 960 gatgtgccaa cagctgccca ggctccgagt cctggaactg tctcacaatc aaattgagga 1020 gctgcccagc ctgcacaggt gtcagaaatt ggaggaaatc ggcctccaac acaaccgcat 1080 ctgggaaatt ggagctgaca ccttcagcca gctgagctcc ctgcaagccc tggatcttag 1140 ctggaacgcc atccggtcca tccaccccga ggccttctcc accctgcact ccctggtcaa 1200 gctggacctg acagacaacc agctgaccac actgcccctg gctggacttg ggggcttgat 1260 gcatctgaag ctcaaaggga accttgctct ctcccaggcc ttctccaagg acagtttccc 1320 aaaactgagg atcctggagg tgccttatgc ctaccagtgc tgtccctatg ggatgtgtgc 1380 cagettette aaggeetetg ggeagtggga ggetgaagae etteacettg atgatgagga 1440 gtcttcaaaa aggcccctgg gcctccttgc cagacaagca gagaaccact atgaccagga 1500 cctggatgag ctccagctgg agatggagga ctcaaagcca caccccagtg tccagtgtag 1560 ccctactcca ggccccttca agccctgtga gtacctcttt gaaagctggg gcatccgcct 1620 ggccgtgtgg gccatcgtgt tgctctccgt gctctgcaat ggactggtgc tgctgaccgt 1680 gttcgctggc gggcctgccc ccctgccccc ggtcaagttt gtggtaggtg cgattgcagg 1740 cgccaacacc ttgactggca tttcctgtgg ccttctagcc tcagtcgatg ccctgacctt 1800 tggtcagttc tctgagtacg gagcccgctg ggagacgggg ctaggctqcc qqqccactqq 1860 cttcctggca gtacttgggt cggaggcatc ggtgctgctg ctcactctgg ccgcagtgca 1920

```
gtgcagcgtc tccgtctcct gtgtccgggc ctatgggaag tccccctccc tgggcaqcqt 1980
tcgagcaggg gtcctaggct gcctggcact ggcagggctg gccgccgcac tgcccctggc 2040
ctcagtggga gaatacgggg cctccccact ctgcctgccc tacgcgccac ctgagggtca 2100
gccagcagcc ctgggcttca ccgtggccct ggtgatgatg aactccttct gtttcctggt 2160
cgtggccggt gcctacatca aactgtactg tgacctgccg cggggcgact ttgaggccgt 2220
gtgggactgc gccatggtga ggcacgtggc ctggctcatc ttcqcaqacq qqctcctcta 2280
ctgtcccgtg gccttcctca gcttcgcctc catgctgggc ctcttccctq tcacqcccqa 2340
ggccgtcaag totgtootgo tggtggtgct gcccctgcct qcctqcctca acccactqct 2400
gtacctgctc ttcaaccccc acttccggga tgaccttcgg cggcttcggc cccgcgcagg 2460
ggactcaggg cccctagcct atgctgcggc cggggagctg qaqaaqaqct cctqtqattc 2520
tacccaggcc ctggtagcct tctctgatgt ggatctcatt ctggaagctt ctgaagctgg 2580
gcggccccct gggctggaga cctatggctt cccctcaqtq accctcatct cctqtcaqca 2640
gccaggggcc cccaggctgg agggcagcca ttgtgtagag ccagagggga accactttgq 2700
gaacccccaa ccctccatgg atggagaact gctgctgagg gcagagggat ctacgccagc 2760
aggtggaggc ttgtcagggg gtggcgcttt caqccctctq qcttqqcctt tqcttcacac 2820
gtgtaaatat ccctccccat tcttctcttc ccctctcttc cctttcctct ctccccctcg 2880
gtgaatgatg gctgcttcta aaacaaatac aaccaaaact cagcagtgtg atctatagca 2940
ggatggccca gtacctggct ccactgatca cctctctcct gtgaccatca ccaacgggtg 3000
ccctcttggc ctggctttcc cttggccttc ctcagcttca ccttgatact gggcctcttc 3060
cttgtcatgt ctgaagctgt ggaccagaga cctggacttt tgtctgctta agggaaatga 3120
gggaagtaaa gacagtgaag gggtggaggg ttgatcaggg cacagtggac agggagacct 3180
cacagagaaa ggcctggaag gtgatttccc gtgtgactca tggataggat acaaaatgtg 3240
ttccatgtac cattaatctt gacatatgcc atgcataaag acttcctatt aaaataagct 3300
ttggaagaga ttacacatga tgtcttttc ttagagattc acagtgcatg ttagtgtaat 3360
aaagagataa gtcctacagt a
                                                                  3381
<210> 6
<213> Homo sapiens
```

```
<211> 940
<212> PRT
```

Met Leu Ser Ala Asp Cys Ser Glu Leu Gly Leu Ser Ala Val Pro Gly

Asp Pro Asp Pro Leu Thr Ala Tyr Leu Asp Leu Ser Met Asn Asn Leu 25

Thr Glu Leu Gln Pro Gly Leu Phe His His Leu Arg Phe Leu Glu Glu

Leu Arg Leu Ser Gly Asn His Leu Ser His Ile Pro Gly Gln Ala Phe

Ser Gly Leu Tyr Ser Leu Lys Ile Leu Met Leu Gln Asn Asn Gln Leu 70 75

Gly Gly Ile Pro Ala Glu Ala Leu Trp Glu Leu Pro Ser Leu Gln Ser 90

Leu Arg Leu Asp Ala Asn Leu Ile Ser Leu Val Pro Glu Arg Ser Phe 105

Glu Gly Leu Ser Ser Leu Arg His Leu Trp Leu Asp Asp Asn Ala Leu 120 125

Thr	Glu 130	Ile	Pro	Val	Arg	Ala 135	Leu	Asn	Asn	Leu	Pro 140	Ala	Leu	Gln	Ala
Met 145	Thr	Leu	Ala	Leu	Asn 150	Arg	Ile	Ser	His	Ile 155	Pro	Asp	Tyr	Ala	Phe 160
Gln	Asn	Leu	Thr	Ser 165	Leu	Val	Val	Leu	His 170	Leu	His	Asn	Asn	Arg 175	Ile
Gln	His	Leu	Gly 180	Thr	His	Ser	Phe	Glu 185	Gly	Leu	His	Asn	Leu 190	Glu	Thr
Leu	Asp	Leu 195	Asn	Tyr	Asn	Lys	Leu 200	Gln	Glu	Phe	Pro	Val 205	Ala	Ile	Arg
Thr	Leu 210	Gly	Arg	Leu	Gln	Glu 215	Leu	Gly	Phe	His	Asn 220	Asn	Asn	Ile	Lys
Ala 225	Ile	Pro	Glu	Lys	Ala 230	Phe	Met	Gly	Asn	Pro 235	Leu	Leu	Gln	Thr	Ile 240
His	Phe	Tyr	Asp	Asn 245	Pro	Ile	Gln	Phe	Val 250	Gly	Arg	Ser	Ala	Phe 255	Gln
Tyr	Leu	Pro	Lys 260	Leu	His	Thr	Leu	Ser 265	Leu	Asn	Gly	Ala	Met 270	Asp	Ile
Gln	Glu	Phe 275	Pro	Asp	Leu	Lys	Gly 280	Thr	Thr	Ser	Leu	Glu 285	Ile	Leu	Thr
Leu	Thr 290	Arg	Ala	Gly	Ile	Arg 295	Leu	Leu	Pro	Ser	Gly 300	Met	Cys	Gln	Gln
Leu 305	Pro	Arg	Leu	Arg	Val 310	Leu	Glu	Leu	Ser	His 315	Asn	Gln	Ile	Glu	Glu 320
Leu	Pro	Ser	Leu	His 325	Arg	Cys	Gln	Lys	Leu 330	Glu	Glu	Ile	Gly	Leu 335	Gln
His	Asn	Arg	Ile 340	Trp	Glu	Ile	Gly	Ala 345	Asp	Thr	Phe	Ser	Gln 350	Leu	Ser
Ser	Leu	Gln 355	Ala	Leu	Asp	Leu	Ser 360	Trp	Asn	Ala	Ile	Arg 365	Ser	Ile	His
Pro	Glu 370	Ala	Phe	Ser	Thr	Leu 375	His	Ser	Leu	Val	Lys 380	Leu	Asp	Leu	Thr
Asp 385	Asn	Gln	Leu	Thr	Thr 390	Leu	Pro	Leu	Ala	Gly 395	Leu	Gly	Gly	Leu	Met 400
His	Leu	Lys	Leu	Lys 405	Gly	Asn	Leu	Ala	Leu 410	Ser	Gln	Ala	Phe	Ser 415	Lys
Asp	Ser	Phe	Pro 420	Lys	Leu	Arg	Ile	Leu 425	Glu	Va]	Pro	_	Ala 430	Tyr	Gln

Cys Cys Pro Tyr Gly Met Cys Ala Ser Phe Phe Lys Ala Ser Gly Gln Trp Glu Ala Glu Asp Leu His Leu Asp Asp Glu Glu Ser Ser Lys Arg 455 Pro Leu Gly Leu Leu Ala Arg Gln Ala Glu Asn His Tyr Asp Gln Asp 470 475 Leu Asp Glu Leu Gln Leu Glu Met Glu Asp Ser Lys Pro His Pro Ser 490 Val Gln Cys Ser Pro Thr Pro Gly Pro Phe Lys Pro Cys Glu Tyr Leu Phe Glu Ser Trp Gly Ile Arg Leu Ala Val Trp Ala Ile Val Leu Leu 520 Ser Val Leu Cys Asn Gly Leu Val Leu Leu Thr Val Phe Ala Gly Gly 535 Pro Ala Pro Leu Pro Pro Val Lys Phe Val Val Gly Ala Ile Ala Gly 550 555 Ala Asn Thr Leu Thr Gly Ile Ser Cys Gly Leu Leu Ala Ser Val Asp Ala Leu Thr Phe Gly Gln Phe Ser Glu Tyr Gly Ala Arg Trp Glu Thr 585 Gly Leu Gly Cys Arg Ala Thr Gly Phe Leu Ala Val Leu Gly Ser Glu 600 Ala Ser Val Leu Leu Thr Leu Ala Ala Val Gln Cys Ser Val Ser 610 615 Val Ser Cys Val Arg Ala Tyr Gly Lys Ser Pro Ser Leu Gly Ser Val Arg Ala Gly Val Leu Gly Cys Leu Ala Leu Ala Gly Leu Ala Ala Ala Leu Pro Leu Ala Ser Val Gly Glu Tyr Gly Ala Ser Pro Leu Cys Leu Pro Tyr Ala Pro Pro Glu Gly Gln Pro Ala Ala Leu Gly Phe Thr Val Ala Leu Val Met Met Asn Ser Phe Cys Phe Leu Val Val Ala Gly Ala Tyr Ile Lys Leu Tyr Cys Asp Leu Pro Arg Gly Asp Phe Glu Ala Val 715 Trp Asp Cys Ala Met Val Arg His Val Ala Trp Leu Ile Phe Ala Asp 725 730

Gly Leu Phe Pro Val Thr Pro Glu Ala Val Lys Ser Val Leu Leu Val 760 765 Val Leu Pro Leu Pro Ala Cys Leu Asn Pro Leu Leu Tyr Leu Leu Phe 775 780 Asn Pro His Phe Arg Asp Asp Leu Arg Arg Leu Arg Pro Arg Ala Gly 785 790 795 Asp Ser Gly Pro Leu Ala Tyr Ala Ala Ala Gly Glu Leu Glu Lys Ser Ser Cys Asp Ser Thr Gln Ala Leu Val Ala Phe Ser Asp Val Asp Leu 820 825 Ile Leu Glu Ala Ser Glu Ala Gly Arg Pro Pro Gly Leu Glu Thr Tyr 835 840 Gly Phe Pro Ser Val Thr Leu Ile Ser Cys Gln Gln Pro Gly Ala Pro 855 860 Arg Leu Glu Gly Ser His Cys Val Glu Pro Glu Gly Asn His Phe Gly 865 870 875 880 Asn Pro Gln Pro Ser Met Asp Gly Glu Leu Leu Arg Ala Glu Gly 885 Ser Thr Pro Ala Gly Gly Gly Leu Ser Gly Gly Gly Ala Phe Ser Pro 905 Leu Ala Trp Pro Leu Leu His Thr Cys Lys Tyr Pro Ser Pro Phe Phe 915 920 925 Ser Ser Pro Leu Phe Pro Phe Leu Ser Pro Pro Arg 935 <210> 7 <211> 2397 <212> DNA <213> Homo sapiens <400> 7 atgctaataa attgtgaagc aaaaggtatc aagatggtat ctgaaataag tgtgccacca 60 tcacgacctt tccaactaag cttattaaat aacggettga cgatgettca cacaaatgac 120 ttttctgggc ttaccaatgc tatttcaata caccttggat ttaacaatat tgcagatatt 180 gagataggtg catttaatgg ccttggcctc ctgaaacaac ttcatatcaa tcacaattct 240 ttagaaattc ttaaagagga tactttccat ggactggaaa acctggaatt cctgcaagca 300 gataacaatt ttatcacagt gattgaacca agtgccttta gcaagctcaa cagactcaaa 360 gtgttaattt taaatgacaa tgctattgag agtcttcctc caaacatctt ccgatttqtt 420 cetttaacce atetagatet tegtggaaat caattacaaa cattgcetta tgttggtttt 480 ctcgaacaca ttggccgaat attggatctt cagttggagg acaacaaatg ggcctgcaat 540 tgtgacttat tgcagttaaa aacttggttg gagaacatgc ctccacagtc tataattggt 600 gatgttgtct gcaacagccc tccatttttt aaaggaagta tactcagtag actaaaqaaq 660

Gly Leu Leu Tyr Cys Pro Val Ala Phe Leu Ser Phe Ala Ser Met Leu

740

```
qaatctattt gccctactcc accagtgtat gaagaacatg aggatccttc aggatcatta 720
catctggcag caacatcttc aataaatgat agtcgcatgt caactaagac cacgtccatt 780
ctaaaactac ccaccaaagc accaggtttg ataccttata ttacaaagcc atccactcaa 840
cttccaggac cttactgccc tattccttgt aactgcaaag tcctatcccc atcaggactt 900
ctaatacatt gtcaggagcg caacattgaa agcttatcag atctgagacc tcctccgcaa 960
aatcctagaa agctcattct agcgggaaat attattcaca gtttaatgaa gtctgatcta 1020
gtggaatatt tcactttgga aatgcttcac ttgggaaaca atcqtattga agttcttgaa 1080
gaaggatcgt ttatgaacct aacgagatta caaaaactct atctaaatgg taaccacctg 1140
accaaattaa gtaaaggcat gttccttggt ctccataatc ttgaatactt atatcttgaa 1200
tacaatgcca ttaaggaaat actgccagga acctttaatc caatgcctaa acttaaagtc 1260
ctgtatttaa ataacaacct cctccaagtt ttaccaccac atattttttc aggggttcct 1320
ctaactaagg taaatcttaa aacaaaccag tttacccatc tacctgtaag taatattttg 1380
gatgatettg atttgetaac ecagattgac ettgaggata acceetggga etgeteetgt 1440
gacctggttg gactgcagca atggatacaa aagttaagca agaacacagt gacagatgac 1500
atcctctgca cttcccccgg gcatctcgac aaaaaggaat tgaaagccct aaatagtgaa 1560
attctctgtc caggtttagt aaataaccca tccatgccaa cacagactag ttaccttatg 1620
gtcaccactc ctgcaacaac aacaaatacg gctgatacta ttttacgatc tcttacggac 1680
gctgtgccac tgtctgttct aatattggga cttctgatta tgttcatcac tattgttttc 1740
tgtgctgcag ggatagtggt tcttgttctt caccgcagga gaagatacaa aaagaaacaa 1800
gtagatgagc aaatgagaga caacagteet gtgeatette agtacageat gtatggeeat 1860
aaaaccactc atcacactac tgaaagaccc tctgcctcac tctatgaaca gcacatggtg 1920
agececatgg tteatgteta tagaagteea teetttggte caaageatet ggaagaggaa 1980
gaagagagga atgagaaaga aggaagtgat gcaaaacatc tccaaagaag tcttttggaa 2040
caggaaaatc attcaccact cacagggtca aatatgaaat acaaaaccac gaaccaatca 2100
acagaatttt tatccttcca agatgccagc tcattgtaca gaaacatttt agaaaaagaa 2160
agggaacttc agcaactggg aatcacagaa tacctaagga aaaacattgc tcaqctccaq 2220
cctgatatgg aggcacatta tcctggagcc cacgaagagc tgaagttaat ggaaacatta 2280
atgtactcac gtccaaggaa ggtattagtg gaacagacaa aaaatgagta ttttgaactt 2340
aaagctaatt tacatgctga acctgactat ttagaagtcc tqqaqcaqca aacatag
```

```
<210> 8
<211> 798
<212> PRT
```

<213> Homo sapiens

<400> 8

Met Leu Ile Asn Cys Glu Ala Lys Gly Ile Lys Met Val Ser Glu Ile 1 5 10 15

Ser Val Pro Pro Ser Arg Pro Phe Gln Leu Ser Leu Leu Asn Asn Gly
20 25 30

Leu Thr Met Leu His Thr Asn Asp Phe Ser Gly Leu Thr Asn Ala Ile
35 40 45

Ser Ile His Leu Gly Phe Asn Asn Ile Ala Asp Ile Glu Ile Gly Ala
50 60

Phe Asn Gly Leu Gly Leu Leu Lys Gln Leu His Ile Asn His Asn Ser 65 70 75 80

Leu Glu Ile Leu Lys Glu Asp Thr Phe His Gly Leu Glu Asn Leu Glu 85 90 95

Phe Leu Gln Ala Asp Asn Asn Phe Ile Thr Val Ile Glu Pro Ser Ala 100 105 110

Phe	Ser	Lys 115	Leu	Asn	Arg	Leu	Lys 120	Val	Leu	Ile	Leu	Asn 125	Asp	Asn	Ala
Ile	Glu 130	Ser	Leu	Pro	Pro	Asn 135	Ile	Phe	Arg	Phe	Val 140	Pro	Leu	Thr	His
Leu 145	Asp	Leu	Arg	Gly	Asn 150	Gln	Leu	Gln	Thr	Leu 155	Pro	Tyr	Val	Gly	Phe 160
Leu	Glu	His	Ile	Gly 165	Arg	Ile	Leu	Asp	Leu 170	Gln	Leu	Glu	Asp	Asn 175	Lys
Trp	Ala	Cys	Asn 180	Cys	Asp	Leu	Leu	Gln 185	Leu	Lys	Thr	Trp	Leu 190	Glu	Asn
Met	Pro	Pro 195	Gln	Ser	Ile	Ile	Gly 200	Asp	Val	Val	Cys	Asn 205	Ser	Pro	Pro
Phe	Phe 210	Lys	Gly	Ser	Ile	Leu 215	Ser	Arg	Leu	Lys	Lys 220	Glu	Ser	Ile	Cys
Pro 225	Thr	Pro	Pro	Val	Tyr 230	Glu	Glu	His	Glu	Asp 235	Pro	Ser	Gly	Ser	Leu 240
His	Leu	Ala	Ala	Thr 245	Ser	Ser	Ile	Asn	Asp 250	Ser	Arg	Met	Ser	Thr 255	Lys
Thr	Thr	Ser	Ile 260	Leu	Lys	Leu	Pro	Thr 265	Lys	Ala	Pro	Gly	Leu 270	Ile	Pro
Tyr	Ile	Thr 275	Lys	Pro	Ser	Thr	Gln 280	Leu	Pro	Gly	Pro	Tyr 285	Cys	Pro	Ile
Pro	Cys 290	Asn	Cys	Lys	Val	Leu 295	Ser	Pro	Ser	Gly	Leu 300	Leu	Ile	His	Cys
Gln 305	Glu	Arg	Asn	Ile	Glu 310	Ser	Leu	Ser	Asp	Leu 315	Arg	Pro	Pro	Pro	Gln 320
Asn	Pro	Arg	Lys	Leu 325	Ile	Leu	Ala	Gly	Asn 330	Ile	Ile	His	Ser	Leu 335	Met
Lys	Ser	Asp	Leu 340	Val	Glu	Tyr	Phe	Thr 345	Leu	Glu	Met	Leu	His 350	Leu	Gly
Asn	Asn	Arg 355	Ile	Glu	Val	Leu	Glu 360	Glu	Gly	Ser	Phe	Met 365	Asn	Leu	Thr
Arg	Leu 370	Gln	Lys	Leu	Tyr	Leu 375	Asn	Gly	Asn	His	Leu 380	Thr	Lys	Leu	Ser
Lys 385	Gly	Met	Phe	Leu	Gly 390	Leu	His	Asn	Leu	Glu 395	Tyr	Leu	Tyr	Leu	Glu 400
Tyr	Asn	Ala	Ile	Lys 405	Glu	Ile	Leu	Pro	Gly 410	Thr	Phe	Asn	Pro	Met 415	Pro

Lys Leu Lys Val Leu Tyr Leu Asn Asn Leu Leu Gln Val Leu Pro 420 425 Pro His Ile Phe Ser Gly Val Pro Leu Thr Lys Val Asn Leu Lys Thr 440 Asn Gln Phe Thr His Leu Pro Val Ser Asn Ile Leu Asp Asp Leu Asp Leu Leu Thr Gln Ile Asp Leu Glu Asp Asn Pro Trp Asp Cys Ser Cys 470 Asp Leu Val Gly Leu Gln Gln Trp Ile Gln Lys Leu Ser Lys Asn Thr 485 490 Val Thr Asp Asp Ile Leu Cys Thr Ser Pro Gly His Leu Asp Lys Lys 500 505 Glu Leu Lys Ala Leu Asn Ser Glu Ile Leu Cys Pro Gly Leu Val Asn 520 Asn Pro Ser Met Pro Thr Gln Thr Ser Tyr Leu Met Val Thr Thr Pro Ala Thr Thr Thr Asn Thr Ala Asp Thr Ile Leu Arg Ser Leu Thr Asp 550 555 Ala Val Pro Leu Ser Val Leu Ile Leu Gly Leu Leu Ile Met Phe Ile 570 Thr Ile Val Phe Cys Ala Ala Gly Ile Val Val Leu Val Leu His Arq Arg Arg Tyr Lys Lys Gln Val Asp Glu Gln Met Arg Asp Asn 600 Ser Pro Val His Leu Gln Tyr Ser Met Tyr Gly His Lys Thr Thr His 610 His Thr Thr Glu Arg Pro Ser Ala Ser Leu Tyr Glu Gln His Met Val 630 Ser Pro Met Val His Val Tyr Arg Ser Pro Ser Phe Gly Pro Lys His Leu Glu Glu Glu Glu Arg Asn Glu Lys Glu Gly Ser Asp Ala Lys 660 His Leu Gln Arg Ser Leu Leu Glu Gln Glu Asn His Ser Pro Leu Thr 680 Gly Ser Asn Met Lys Tyr Lys Thr Thr Asn Gln Ser Thr Glu Phe Leu 690 695 Ser Phe Gln Asp Ala Ser Ser Leu Tyr Arg Asn Ile Leu Glu Lys Glu 705 710 715 720

```
Arg Glu Leu Gln Gln Leu Gly Ile Thr Glu Tyr Leu Arg Lys Asn Ile
725 730 735
```

Ala Gln Leu Gln Pro Asp Met Glu Ala His Tyr Pro Gly Ala His Glu
740 745 750

Glu Leu Lys Leu Met Glu Thr Leu Met Tyr Ser Arg Pro Arg Lys Val
755 760 765

Leu Val Glu Gln Thr Lys Asn Glu Tyr Phe Glu Leu Lys Ala Asn Leu 770 780

His Ala Glu Pro Asp Tyr Leu Glu Val Leu Glu Gln Gln Thr
785 790 795

```
<210> 9
<211> 3825
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (2)..(7)
<223> Wherein n is a or t or c or g.
```

<220>
<221> misc\_feature

<222> (10)..(12)

<223> Wherein n is a or t or c or g.

<220> <221> mis

<221> misc\_feature

<222> (17)

<223> Wherein n is a or t or c or g.

<220>

<221> misc\_feature

<222> (19)

<223> Wherein n is a or t or c or g.

<220>

<221> misc\_feature

<222> (22)

<223> Wherein n is a or t or c or g.

<220>

<221> misc\_feature

<222> (25)

<223> Wherein n is a or t or c or g.

<220>

<221> misc\_feature

<222> (28)..(30)

<223> Wherein n is a or t or c or g.

aaacctacag catattette acgeagagat ttteatetat tataetttat eaaagattgg 3300 ceatgtteea ettggaaatg geatgeaaaa geeateatag agaaacetge gtaaeteeat 3360

ctgacaaatt caaaagagag agagagatct tgagagagaa atgctgttcg ttcaaaagtg 3420 gagttgtttt aacagatgcc aattacggtg tacagtttaa cagagttttc tgttgcatta 3480 ggataaacat taattggagt gcagctaaca tgagtatcat cagactagta tcaagtgttc 3540 taaaatgaaa tatgagaaga tcctgtcaca attcttagat ctggtgtcca gcatggatga 3600 aacctttgag tttggtcct aaatttgcat gaaagcacaa ggtaaatatt catttgcttc 3660 aggagtttca tgttggatct gtcattatca aaagtgatca gcaatgaaga actggtcgga 3720 caaaatttaa cgttgatga ttcatttgaa taaaaaggaa cttgg 3825

<210> 10

<211> 865

<212> PRT

<213> Homo sapiens

<400> 10

Met Ala Leu Val Leu Gly Ser Leu Leu Leu Leu Gly Leu Cys Gly Asn 1 5 10 15

Ser Phe Ser Gly Gly Gln Pro Ser Ser Thr Asp Ala Pro Lys Ala Trp
20 25 30

Asn Tyr Glu Leu Pro Ala Thr Asn Tyr Glu Thr Gln Asp Ser His Lys
35 40 45

Ala Gly Pro Ile Gly Ile Leu Phe Glu Leu Val His Ile Phe Leu Tyr 50 55 60

Val Val Gln Pro Arg Asp Phe Pro Glu Asp Thr Leu Arg Lys Phe Leu 65 70 75 80

Gln Lys Ala Tyr Glu Ser Lys Ile Asp Tyr Asp Lys Pro Glu Thr Val 85 90 95

Ile Leu Gly Leu Lys Ile Val Tyr Tyr Glu Ala Gly Ile Ile Leu Cys 100 105 110

Cys Val Leu Gly Leu Leu Phe Ile Ile Leu Met Pro Leu Val Gly Tyr 115 120 125

Phe Phe Cys Met Cys Arg Cys Cys Asn Lys Cys Gly Glu Met His 130 135 140

Gln Arg Gln Lys Glu Asn Gly Pro Phe Leu Arg Lys Cys Phe Ala Ile 145 150 155 160

Ser Leu Leu Val Ile Cys Ile Ile Ile Ser Ile Gly Ile Phe Tyr Gly
165 170 175

Phe Val Ala Asn His Gln Val Arg Thr Arg Ile Lys Arg Ser Arg Lys 180 185 190

Leu Ala Asp Ser Asn Phe Lys Asp Leu Arg Thr Leu Leu Asn Glu Thr
195 200 205

Pro Glu Gln Ile Lys Tyr Ile Leu Ala Gln Tyr Asn Thr Thr Lys Asp 210 215 220

Lys 225	Ala	Phe	Thr	Asp	Leu 230	Asn	Ser	Ile	Asn	Ser 235	Val	Leu	Gly	Gly	Gly 240
Ile	Leu	Asp	Arg	Leu 245	Arg	Pro	Asn	Ile	Ile 250	Pro	Val	Leu	Asp	Glu 255	Ile
Lys	Ser	Met	Ala 260	Thr	Ala	Ile	Lys	Glu 265	Thr	Lys	Glu	Ala	Leu 270	Glu	Asn
Met	Asn	Ser 275	Thr	Leu	Lys	Ser	Leu 280	His	Gln	Gln	Ser	Thr 285	Gln	Leu	Ser
Ser	Ser 290	Leu	Thr	Ser	Val	Lys 295	Thr	Ser	Leu	Arg	Ser 300	Ser	Leu	Asn	Asp
Pro 305	Leu	Cys	Leu	Val	His 310	Pro	Ser	Ser	Glu	Thr 315	Cys	Asn	Ser	Ile	Arg 320
Leu	Ser	Leu	Ser	Gln 325	Leu	Asn	Ser	Asn	Pro 330	Glu	Leu	Arg	Gln	Leu 335	Pro
Pro	Val	Asp	Ala 340	Glu	Leu	Asp	Asn	Val 345	Asn	Asn	Val	Leu	Arg 350	Thr	Asp
Leu	Asp	Gly 355	Leu	Val	Gln	Gln	Gly 360	Tyr	Gln	Ser	Leu	Asn 365	Asp	Ile	Pro
Asp	Arg 370	Val	Gln	Arg	Gln	Thr 375	Thr	Thr	Val	Val	Ala 380	Gly	Ile	Lys	Arg
Val 385	Leu	Asn	Ser	Ile	Gly 390	Ser	Asp	Ile	Asp	Asn 395	Val	Thr	Gln	Arg	Leu 400
Pro	Ile	Gln	Asp	Ile 405	Leu	Ser	Ala	Phe	Ser 410	Val	Tyr	Val	Asn	Asn 415	Thr
Glu	Ser	Tyr	Ile 420	His	Arg	Asn	Leu	Pro 425	Thr	Leu	Glu	Glu	Tyr 430	Asp	Ser
Tyr	Trp	Trp 435	Leu	Gly	Gly	Leu	Val 440	Ile	Cys	Ser	Leu	Leu 445	Thr	Leu	Ile
Val	Ile 450	Phe	Tyr	Tyr	Leu	Gly 455	Leu	Leu	Cys	Gly	Val 460	Cys	Gly	Tyr	Asp
Arg 465	His	Ala	Thr	Pro	Thr 470	Thr	Arg	Gly	Cys	Val 475	Ser	Asn	Thr	Gly	Gly 480
Val	Phe	Leu	Met	Val 485	Gly	Val	Gly	Leu	Ser 490	Phe	Leu	Phe	Cys	Trp 495	Ile
Leu	Met	Ile	Ile 500	Val	Val	Leu	Thr	Phe 505	Val	Phe	Gly	Ala	Asn 510	Val	Glu
Lys	Leu	Ile 515	Cys	Glu	Pro	Tyr	Thr 520	Ser	Lys	Glu	Leu	Phe 525	Arg	Val	Leu

Asp Thr Pro Tyr Leu Leu Asn Glu Asp Trp Glu Tyr Tyr Leu Ser Gly 535 Lys Leu Phe Asn Lys Ser Lys Met Lys Leu Thr Phe Glu Gln Val Tyr 550 Ser Asp Cys Lys Asn Arg Gly Thr Tyr Gly Thr Leu His Leu Gln 565 Asn Ser Phe Asn Ile Ser Glu His Leu Asn Ile Asn Glu His Thr Gly 585 Ser Ile Ser Ser Glu Leu Glu Ser Leu Lys Val Asn Leu Asn Ile Phe Leu Leu Gly Ala Ala Gly Arg Lys Asn Leu Gln Asp Phe Ala Ala Cys 615 Gly Ile Asp Arg Met Asn Tyr Asp Ser Tyr Leu Ala Gln Thr Gly Lys Ser Pro Ala Gly Val Asn Leu Leu Ser Phe Ala Tyr Asp Leu Glu Ala 645 650 Lys Ala Asn Ser Leu Pro Pro Gly Asn Leu Arg Asn Ser Leu Lys Arg 665 Asp Ala Gln Thr Ile Lys Thr Ile His Gln Gln Arg Val Leu Pro Ile Glu Gln Ser Leu Ser Thr Leu Tyr Gln Ser Val Lys Ile Leu Gln Arg Thr Gly Asn Gly Leu Leu Glu Arg Val Thr Arg Ile Leu Ala Ser Leu 710 715 Asp Phe Ala Gln Asn Phe Ile Thr Asn Asn Thr Ser Ser Val Ile Ile 725 Glu Glu Thr Lys Lys Tyr Gly Arg Thr Ile Ile Gly Tyr Phe Glu His Tyr Leu Gln Trp Ile Glu Phe Ser Ile Ser Glu Lys Val Ala Ser Cys 760 Lys Pro Val Ala Thr Ala Leu Asp Thr Ala Val Asp Val Phe Leu Cys Ser Tyr Ile Ile Asp Pro Leu Asn Leu Phe Trp Phe Gly Ile Gly Lys 790 795 Ala Thr Val Phe Leu Leu Pro Ala Leu Ile Phe Ala Val Lys Leu Ala 805 Lys Tyr Tyr Arg Arg Met Asp Ser Glu Asp Val Tyr Asp Asp Val Glu 820 825 830

```
Thr Ile Pro Met Lys Asn Met Glu Asn Gly Asn Asn Gly Tyr His Lys
                            840
Asp His Val Tyr Gly Ile His Asn Pro Val Met Thr Ser Pro Ser Gln
His
865
<210> 11
<211> 1807
<212> DNA
<213> Homo sapiens
<400> 11
gcacgaggga agagggtgat ccgacccggg gaaggtcgct gggcagggcg agttgggaaa 60
geggeageee eegeegeeee egeageeeet teteeteett teteeeaegt eetatetgee 120
tetegetgga ggeeaggeeg tgeageateg aagacaggag gaactggage eteattqqee 180
ggcccggggc gccggcctcg ggcttaaata ggagctccgg gctctggctg ggacccgacc 240
getgeeggee gegeteeege tgeteetgee gggtgatgga aaaccccage eeggeegeeg 300
ccctgggcaa ggccctctgc gctctcctcc tggccactct cggcgccgcc ggccagcctc 360
ttgggggaga gtccatctgt tccgccagag ccccggccaa atacagcatc accttcacgg 420
gcaagtggag ccagacggcc ttccccaagc agtaccccct gttccgcccc cctgcgcagt 480
ggtcttcgct gctgggggcc gcgcatagct ccgactacag catgtggagg aagaaccagt 540
acgtcagtaa cgggctgcgc gactttgcgg agcgcggcga ggcctgggcg ctgatgaagg 600
agategagge ggegggggag gegetgeaga gegtgeaege ggtgtttteg gegeeegeeg 660
tccccagcgg caccgggcag acgtcggcgg agctggaggt gcagcgcagg cactcgctgg 720
tetegtttgt ggtgegeate gtgeeeagee eegactggtt egtgggegtg gaeageetgg 780
acctgtgcga cggggaccgt tggcgggaac aggcggcgct ggacctgtac ccctacqacq 840
ccgggacgga cagcggcttc accttctcct ccccaactt cgccaccatc ccqcaqqaca 900
cggtgaccga gataacgtcc tcctctccca gccacccggc caactccttc tactacccqc 960
ggctgaaggc cctgcctccc atcgccaggg tgacactggt gcggctgcga caqaqcccca 1020
gggccttcat ccctcccgcc ccagtcctgc ccagcaggga caatgagatt gtagacagcg 1080
cctcagttcc agaaacgccg ctggactgcg aggtctccct gtggtcgtcc tggggactqt 1140
geggaggeea etgtgggagg etegggaeea agageaggae tegetaegte egggteeage 1200
ccgccaacaa cgggagcccc tgccccgagc tcgaagaaga ggctgagtgc gtccctgata 1260
actgcgtcta agaccagagc cccgcagccc ctggggcccc cggagccatg gggtgtcgqg 1320
ggeteetgtg caggeteatg etgeaggegg eegaggeaca gggggttteg egetgeteet 1380
gaccgcggtg aggccgccc gaccatctct gcactgaagg gccctctggt ggccggcacg 1440
ggcattggga aacagcctcc tcctttccca accttgcttc ttaggggccc ccgtgtcccg 1500
tetgetetea geeteeteet eetgeaggat aaagteatee eeaaggetee agetaeteta 1560
aattatggtc tccttataag ttattgctgc tccaggagat tgtccttcat cgtccagggg 1620
cctggctccc acgtggttgc agatacctca gacctggtgc tctaggctgt gctgagccca 1680
ctctcccgag ggcgcatcca agcgggggcc acttgagaag tgaataaatg gggcggtttc 1740
ggaagcgtca gtgtttccat gttatggatc tctctgcgtt tgaataaaga ctatctctgt 1800
tgctcac
                                                                  1807
<210> 12
<211> 331
<212> PRT
<213> Homo sapiens
<400> 12
```

Met Glu Asn Pro Ser Pro Ala Ala Ala Leu Gly Lys Ala Leu Cys Ala

- Leu Leu Leu Ala Thr Leu Gly Ala Ala Gly Gln Pro Leu Gly Gly Glu 20 25 30
- Ser Ile Cys Ser Ala Arg Ala Pro Ala Lys Tyr Ser Ile Thr Phe Thr 35 40 45
- Gly Lys Trp Ser Gln Thr Ala Phe Pro Lys Gln Tyr Pro Leu Phe Arg
  50 55 60
- Pro Pro Ala Gln Trp Ser Ser Leu Leu Gly Ala Ala His Ser Ser Asp 65 70 75 80
- Tyr Ser Met Trp Arg Lys Asn Gln Tyr Val Ser Asn Gly Leu Arg Asp 85 90 95
- Phe Ala Glu Arg Gly Glu Ala Trp Ala Leu Met Lys Glu Ile Glu Ala 100 105 110
- Ala Gly Glu Ala Leu Gln Ser Val His Ala Val Phe Ser Ala Pro Ala 115 120 125
- Val Pro Ser Gly Thr Gly Gln Thr Ser Ala Glu Leu Glu Val Gln Arg 130 135 140
- Arg His Ser Leu Val Ser Phe Val Val Arg Ile Val Pro Ser Pro Asp 145 150 155 160
- Trp Phe Val Gly Val Asp Ser Leu Asp Leu Cys Asp Gly Asp Arg Trp
  165 170 175
- Arg Glu Gln Ala Ala Leu Asp Leu Tyr Pro Tyr Asp Ala Gly Thr Asp 180 185 190
- Ser Gly Phe Thr Phe Ser Ser Pro Asn Phe Ala Thr Ile Pro Gln Asp 195 200 205
- Thr Val Thr Glu Ile Thr Ser Ser Ser Pro Ser His Pro Ala Asn Ser 210 215 220
- Phe Tyr Tyr Pro Arg Leu Lys Ala Leu Pro Pro Ile Ala Arg Val Thr 225 230 235 240
- Leu Val Arg Leu Arg Gln Ser Pro Arg Ala Phe Ile Pro Pro Ala Pro
  245 250 255
- Val Leu Pro Ser Arg Asp Asn Glu Ile Val Asp Ser Ala Ser Val Pro 260 265 270
- Glu Thr Pro Leu Asp Cys Glu Val Ser Leu Trp Ser Ser Trp Gly Leu 275 280 285
- Cys Gly Gly His Cys Gly Arg Leu Gly Thr Lys Ser Arg Thr Arg Tyr
  290 295 300
- Val Arg Val Gln Pro Ala Asn Asn Gly Ser Pro Cys Pro Glu Leu Glu

Glu Glu Ala Glu Cys Val Pro Asp Asn Cys Val 325 330

<210> 13

<211> 898

<212> PRT

<213> Rattus norvegicus

<400> 13

Met Ala Val Arg Pro Gly Leu Trp Pro Val Leu Leu Gly Ile Val Leu

1 5 10 15

Ala Ala Trp Leu Arg Gly Ser Gly Ala Gln Gln Ser Ala Thr Val Ala 20 25 30

Asn Pro Val Pro Gly Ala Asn Pro Asp Leu Leu Pro His Phe Leu Val
35 40 45

Glu Pro Glu Asp Val Tyr Ile Val Lys Asn Lys Pro Val Leu Leu Val
50 55 60

Cys Lys Ala Val Pro Ala Thr Gln Ile Phe Phe Lys Cys Asn Gly Glu 65 70 75 80

Trp Val Arg Gln Val Asp His Val Ile Glu Arg Ser Thr Asp Ser Ser 85 90 95

Ser Gly Leu Pro Thr Met Glu Val Arg Ile Asn Val Ser Arg Gln Gln 100 105 110

Val Glu Lys Val Phe Gly Leu Glu Glu Tyr Trp Cys Gln Cys Val Ala 115 120 125

Trp Ser Ser Ser Gly Thr Thr Lys Ser Gln Lys Ala Tyr Ile Arg Ile 130 135 140

Ser Leu Glu Gln Gly Ile Val Leu Pro Cys Arg Pro Pro Glu Gly Ile 165 170 175

Pro Pro Ala Glu Val Glu Trp Leu Arg Asn Glu Asp Leu Val Asp Pro 180 185 190

Ser Leu Asp Pro Asn Val Tyr Ile Thr Arg Glu His Ser Leu Val Val 195 200 205

Arg Gln Ala Arg Leu Ala Asp Thr Ala Asn Tyr Thr Cys Val Ala Lys 210 215 220

Asn Ile Val Ala Arg Arg Ser Thr Ser Ala Ala Val Ile Val Tyr 225 230 235 240

Val Asn Gly Gly Trp Ser Thr Trp Thr Glu Trp Ser Val Cys Ser Ala Ser Cys Gly Arg Gly Trp Gln Lys Arg Ser Arg Ser Cys Thr Asn Pro Ala Pro Leu Asn Gly Gly Ala Phe Cys Glu Gly Gln Asn Val Gln Lys 280 Thr Ala Cys Ala Thr Leu Cys Pro Val Asp Gly Ser Trp Ser Ser Trp 290 Ser Lys Trp Ser Ala Cys Gly Leu Asp Cys Thr His Trp Arg Ser Arg Glu Cys Ser Asp Pro Ala Pro Arg Asn Gly Glu Glu Cys Arg Gly 330 Ala Asp Leu Asp Thr Arg Asn Cys Thr Ser Asp Leu Cys Leu His Thr 345 Ala Ser Cys Pro Glu Asp Val Ala Leu Tyr Ile Gly Leu Val Ala Val 360 Ala Val Cys Leu Phe Leu Leu Leu Ala Leu Gly Leu Ile Tyr Cys 370 Arg Lys Lys Glu Gly Leu Asp Ser Asp Val Ala Asp Ser Ser Ile Leu Thr Ser Gly Phe Gln Pro Val Ser Ile Lys Pro Ser Lys Ala Asp Asn 410 Pro His Leu Leu Thr Ile Gln Pro Asp Leu Ser Thr Thr Thr Thr 420 425 Tyr Gln Gly Ser Leu Cys Ser Arg Gln Asp Gly Pro Ser Pro Lys Phe 440 Gln Leu Ser Asn Gly His Leu Leu Ser Pro Leu Gly Ser Gly Arg His 450 Thr Leu His His Ser Ser Pro Thr Ser Glu Ala Glu Asp Phe Val Ser Arg Leu Ser Thr Gln Asn Tyr Phe Arg Ser Leu Pro Arg Gly Thr Ser 490 Asn Met Ala Tyr Gly Thr Phe Asn Phe Leu Gly Gly Arg Leu Met Ile 500 505 Pro Asn Thr Gly Ile Ser Leu Leu Ile Pro Pro Asp Ala Ile Pro Arg 520 Gly Lys Ile Tyr Glu Ile Tyr Leu Thr Leu His Lys Pro Glu Asp Val

Arg Leu Pro Leu Ala Gly Cys Gln Thr Leu Leu Ser Pro Val Val Ser 545 Cys Gly Pro Pro Gly Val Leu Leu Thr Arg Pro Val Ile Leu Ala Met 565 570 Asp His Cys Gly Glu Pro Ser Pro Asp Ser Trp Ser Leu Arg Leu Lys 585 Lys Gln Ser Cys Glu Gly Ser Trp Glu Asp Val Leu His Leu Gly Glu 600 Glu Ser Pro Ser His Leu Tyr Tyr Cys Gln Leu Glu Ala Gly Ala Cys Tyr Val Phe Thr Glu Gln Leu Gly Arg Phe Ala Leu Val Gly Glu Ala Leu Ser Val Ala Ala Thr Lys Arg Leu Arg Leu Leu Phe Ala Pro Val Ala Cys Thr Ser Leu Glu Tyr Asn Ile Arg Val Tyr Cys Leu His 665 Asp Thr His Asp Ala Leu Lys Glu Val Val Gln Leu Glu Lys Gln Leu 675 680 Gly Gly Gln Leu Ile Gln Glu Pro Arg Val Leu His Phe Lys Asp Ser Tyr His Asn Leu Arg Leu Ser Ile His Asp Val Pro Ser Ser Leu Trp 710 715 Lys Ser Lys Leu Leu Val Ser Tyr Gln Glu Ile Pro Phe Tyr His Ile 725 730 Trp Asn Gly Thr Gln Gln Tyr Leu His Cys Thr Phe Thr Leu Glu Arg 745 Ile Asn Ala Ser Thr Ser Asp Leu Ala Cys Lys Val Trp Val Trp Gln Val Glu Gly Asp Gly Gln Ser Phe Asn Ile Asn Phe Asn Ile Thr Lys 775 Asp Thr Arg Phe Ala Glu Leu Leu Ala Leu Glu Ser Glu Gly Gly Val 790 795 Pro Ala Leu Val Gly Pro Ser Ala Phe Lys Ile Pro Phe Leu Ile Arg 805 Gln Lys Ile Ile Ala Ser Leu Asp Pro Pro Cys Ser Arg Gly Ala Asp 825 Trp Arg Thr Leu Ala Gln Lys Leu His Leu Asp Ser His Leu Ser Phe

Phe Ala Ser Lys Pro Ser Pro Thr Ala Met Ile Leu Asn Leu Trp Glu 850 855 860

Ala Arg His Phe Pro Asn Gly Asn Leu Gly Gln Leu Ala Ala Val 865 870 875 880

Ala Gly Leu Gly Gln Pro Asp Ala Gly Leu Phe Thr Val Ser Glu Ala 885 890 895

Glu Cys

<210> 14

<211> 544

<212> PRT

<213> Homo sapiens

<400> 14

Ala Arg Gly Asp Val Ala Leu Tyr Val Gly Leu Ile Ala Val Ala Val 1 5 10 15

Cys Leu Val Leu Leu Leu Val Leu Ile Leu Val Tyr Cys Arg Lys
20 25 30

Lys Glu Gly Leu Asp Ser Asp Val Ala Asp Ser Ser Ile Leu Thr Ser 35 40 45

Gly Phe Gln Pro Val Ser Ile Lys Pro Ser Lys Ala Asp Asn Pro His 50 55 60

Leu Leu Thr Ile Gln Pro Asp Leu Ser Thr Thr Thr Thr Thr Tyr Gln 65 70 75 80

Gly Ser Leu Cys Pro Arg Gln Asp Gly Pro Ser Pro Lys Phe Gln Leu 85 90 95

Thr Asn Gly His Leu Leu Ser Pro Leu Gly Gly Gly Arg His Thr Leu
100 105 110

His His Ser Ser Pro Thr Ser Glu Ala Glu Glu Phe Val Ser Arg Leu 115 120 125

Ser Thr Gln Asn Tyr Phe Arg Ser Leu Pro Arg Gly Thr Ser Asn Met 130 135 140

Thr Tyr Gly Thr Phe Asn Phe Leu Gly Gly Arg Leu Met Ile Pro Asn 145 150 155 160

Thr Gly Ile Ser Leu Leu Ile Pro Pro Asp Ala Ile Pro Arg Gly Lys 165 170 175

Ile Tyr Glu Ile Tyr Leu Thr Leu His Lys Pro Glu Asp Val Arg Leu
180 185 190

Pro Leu Ala Gly Cys Gln Thr Leu Leu Ser Pro Ile Val Ser Cys Gly
195 200 205

Pro Pro Gly Val Leu Leu Thr Arg Pro Val Ile Leu Ala Met Asp His Cys Gly Glu Pro Ser Pro Asp Ser Trp Ser Leu Arg Leu Lys Lys Gln 230 Ser Cys Glu Gly Ser Trp Glu Asp Val Leu His Leu Gly Glu Glu Ala Pro Ser His Leu Tyr Tyr Cys Gln Leu Glu Ala Ser Ala Cys Tyr Val 265 Phe Thr Glu Gln Leu Gly Arg Phe Ala Leu Val Gly Glu Ala Leu Ser 280 Val Ala Ala Ala Lys Arg Leu Lys Leu Leu Phe Ala Pro Val Ala Cys Thr Ser Leu Glu Tyr Asn Ile Arg Val Tyr Cys Leu His Asp Thr 310 His Asp Ala Leu Lys Glu Val Val Gln Leu Glu Lys Gln Leu Gly Gly 325 Gln Leu Ile Gln Glu Pro Arg Val Leu His Phe Lys Asp Ser Tyr His 345 Asn Leu Arg Leu Ser Ile His Asp Val Pro Ser Ser Leu Trp Lys Ser 360 Lys Leu Leu Val Ser Tyr Gln Glu Ile Pro Phe Tyr His Ile Trp Asn 375 Gly Thr Gln Arg Tyr Leu His Cys Thr Phe Thr Leu Glu Arg Val Ser 390 395 Pro Ser Thr Ser Asp Leu Ala Cys Lys Leu Trp Val Trp Gln Val Glu 405 Gly Asp Gly Gln Ser Phe Ser Ile Asn Phe Asn Ile Thr Lys Asp Thr Arg Phe Ala Glu Leu Leu Ala Leu Glu Ser Glu Ala Gly Val Pro Ala 435 440 Leu Val Gly Pro Ser Ala Phe Lys Ile Pro Phe Leu Ile Arg Gln Lys 450 Ile Ile Ser Ser Leu Asp Pro Pro Cys Arg Arg Gly Ala Asp Trp Arg 470 475 Thr Leu Ala Gln Lys Leu His Leu Asp Ser His Leu Ser Phe Phe Ala 485 Ser Lys Pro Ser Pro Thr Ala Met Ile Leu Asn Leu Trp Glu Ala Arq 500 505 510

- His Phe Pro Asn Gly Asn Leu Ser Gln Leu Ala Ala Ala Val Ala Gly 515 520 525
- Leu Gly Gln Pro Asp Ala Gly Leu Phe Thr Val Ser Glu Ala Glu Cys 530 535 540
- <210> 15
- <211> 931
- <212> PRT
- <213> Caenorhabditis elegans
- <400> 15
- Met Arg Lys Gly Leu Arg Ala Thr Ala Ala Arg Cys Gly Leu Gly Leu 1 5 10 15
- Gly Tyr Leu Leu Gln Met Leu Val Leu Pro Ala Leu Ala Leu Leu Ser 20 25 30
- Ala Ser Gly Thr Gly Ser Ala Ala Gln Asp Asp Glu Phe Phe His Glu 35 40 45
- Leu Pro Glu Thr Phe Pro Ser Asp Pro Pro Glu Pro Leu Pro His Phe 50 55 60
- Leu Ile Glu Pro Glu Glu Ala Tyr Ile Val Lys Asn Lys Pro Val Asn 65 70 75 80
- Leu Tyr Cys Lys Ala Ser Pro Ala Thr Gln Ile Tyr Phe Lys Cys Asn 85 90 95
- Ser Glu Trp Val His Gln Lys Asp His Val Val Asp Glu Arg Val Asp 100 105 110
- Glu Thr Ser Gly Leu Ile Val Arg Glu Val Ser Ile Glu Ile Ser Arg 115 120 125
- Gln Gln Val Glu Glu Leu Phe Gly Pro Glu Asp Tyr Trp Cys Gln Cys 130 135 140
- Val Ala Trp Ser Ser Ala Gly Thr Thr Lys Ser Arg Lys Ala Tyr Val 145 150 155 160
- Arg Ile Ala Tyr Leu Arg Lys Thr Phe Glu Gln Glu Pro Leu Gly Lys
  165 170 175
- Glu Val Ser Leu Glu Gln Glu Val Leu Leu Gln Cys Arg Pro Pro Glu 180 185 190
- Gly Ile Pro Val Ala Glu Val Glu Trp Leu Lys Asn Glu Asp Ile Ile 195 200 205
- Asp Pro Ala Glu Asp Arg Asn Phe Tyr Ile Thr Ile Asp His Asn Leu

	210					215					220			
Ile 225	Ile	Lys	Gln	Ala	Arg 230	Leu	Ser	Asp	Thr	Ala 235	Asn	Tyr	Thr	Cys
Ala	Lys	Asn	Ile	Val 245	Ala	Lys	Arg	Lys	Ser 250	Thr	Thr	Ala	Thr	Val 255
Val	Tyr	Val	Asn 260	Gly	Gly	Trp	Ser	Thr 265	Trp	Thr	Glu	Trp	Ser 270	Val
Asn	Ser	Arg 275	Cys	Gly	Arg	Gly	Tyr 280	Gln	Lys	Arg	Thr	Arg 285	Thr	Cys

Val 240

Ile

Cys

Thr

Asn Pro Ala Pro Leu Asn Gly Gly Ala Phe Cys Glu Gly Gln Ser Val 290 295 300

Gln Lys Ile Ala Cys Thr Thr Leu Cys Pro Val Asp Gly Arg Trp Thr 305 310 315 320

Ser Trp Ser Lys Trp Ser Thr Cys Gly Thr Glu Cys Thr His Trp Arg 325 330 335

Arg Arg Glu Cys Thr Ala Pro Ala Pro Lys Asn Gly Gly Lys Asp Cys 340 345 350

Asp Gly Leu Val Leu Gln Ser Lys Asn Cys Thr Asp Gly Leu Cys Met 355 360 365

Gln Ala Ala Pro Asp Ser Asp Asp Val Ala Leu Tyr Val Gly Ile Val 370 375 380

Ile Ala Val Thr Val Cys Leu Ala Ile Thr Val Val Val Ala Leu Phe 385 390 395 400

Val Tyr Arg Lys Asn His Arg Asp Phe Glu Ser Asp Ile Ile Asp Ser 405 410 415

Ser Ala Leu Asn Gly Gly Phe Gln Pro Val Asn Ile Lys Ala Ala Arg 420 425 430

Gln Asp Leu Leu Ala Val Pro Pro Asp Leu Thr Ser Ala Ala Met 435 440 445

Tyr Arg Gly Pro Val Tyr Ala Leu His Asp Val Ser Asp Lys Ile Pro 450 455 460

Met Thr Asn Ser Pro Ile Leu Asp Pro Leu Pro Asn Leu Lys Ile Lys 465 470 475 480

Val Tyr Asn Ser Ser Gly Ala Val Thr Pro Gln Asp Asp Leu Ala Glu 485 490 495

Phe Ser Ser Lys Leu Ser Pro Gln Met Thr Gln Ser Leu Leu Glu Asn 500 505 510

Glu Ala Leu Asn Leu Lys Asn Gln Ser Leu Ala Arg Gln Thr Asp Pro

Ser	Cys 530	Thr	Ala	Phe	Gly	Thr 535	Phe	Asn	Ser	Leu	Gly 540	Gly	His	Leu	Ile
Ile 545	Pro	Asn	Ser	Gly	Val 550	Ser	Leu	Leu	Ile	Pro 555	Ala	Gly	Ala	Ile	Pro 560
Gln	Gly	Arg	Val	Tyr 565	Glu	Met	Tyr	Val	Thr 570	Val	His	Arg	Lys	Glu 575	Ası
Met	Arg	Pro	Pro 580	Met	Glu	Asp	Ser	Gln 585	Thr	Leu	Leu	Thr	Pro 590	Val	Va.
Ser	Cys	Gly 595	Pro	Pro	Gly	Ala	Leu 600	Leu	Thr	Arg	Pro	Val 605	Ile	Leu	Thi
Leu	His 610	His	Cys	Ala	Asp	Pro 615	Ser	Thr	Glu	Asp	Trp 620	Lys	Ile	Gln	Leı
Lys 625	Asn	Gln	Ala	Val	Gln 630	Gly	Gln	Trp	Glu	Asp 635	Val	Val	Val	Val	Gl <sub>y</sub> 640
Glu	Glu	Asn	Phe	Thr 645	Thr	Pro	Cys	Tyr	Ile 650	Gln	Leu	Asp	Ala	Glu 655	Ala
Cys	His	Ile	Leu 660	Thr	Glu	Asn	Leu	Ser 665	Thr	Tyr	Ala	Leu	Val 670	Gly	Glr
Ser	Thr	Thr 675	Lys	Ala	Ala	Ala	Lys 680	Arg	Leu	Lys	Leu	Ala 685	Ile	Phe	Gly
Pro	Leu 690	Cys	Cys	Ser	Ser	Leu 695	Glu	Tyr	Ser	Ile	Arg 700	Val	Tyr	Cys	Leu
Asp 705	Asp	Thr	Gln	Asp	Ala 710	Leu	Lys	Glu	Val	Leu 715	Gln	Leu	Glu	Arg	Glr 720
Met	Gly	Gly	Gln	Leu 725	Leu	Glu	Glu	Pro	Lys 730	Ala	Leu	His	Phe	Lys 735	Gly
Ser	Ile	His	Asn 740	Leu	Arg	Leu	Ser	Ile 745	His	Asp	Ile	Ala	His 750	Ser	Leu
Trp	Lys	Ser 755	Lys	Leu	Leu	Ala	Lys 760	Tyr	Gln	Glu	Ile	Pro 765	Phe	Tyr	His
Ile	Trp 770	Ser	Gly	Ser	Gln	Arg 775	Asn	Leu	His	Cys	Thr 780	Phe	Thr	Leu	Glu
Arg 785	Leu	Ser	Leu	Asn	Thr 790	Val	Glu	Leu	Val	Cys 795	Lys	Leu	Cys	Val	Arg 800
Gln	Val	Glu	Gly	Glu 805	Gly	Gln	Ile	Phe	Gln 810	Leu	Asn	Cys	Thr	Val 815	Ser
Glu	Glu	Pro	Thr	Glv	Tle	Agn	Len	Pro	Len	Len	Aen	Dro	Δl =	Ser	Thy

820 825 8	330
-----------	-----

Ile Thr Thr Val Thr Gly Pro Ser Ala Phe Ser Ile Pro Leu Pro Ile 835 840 845

Arg Gln Lys Leu Cys Ser Ser Leu Asp Ala Pro Gln Thr Arg Gly His 850 860

Asp Trp Arg Met Leu Ala His Lys Leu Asn Leu Asp Arg Tyr Leu Asn 865 870 875 880

Tyr Phe Ala Thr Lys Ser Ser Pro Thr Gly Val Ile Leu Asp Leu Trp 885 890 895

Glu Ala Gln Asn Phe Pro Asp Gly Asn Leu Ser Met Leu Ala Ala Val $900 \hspace{1.5cm} 905 \hspace{1.5cm} 910$ 

Leu Glu Glu Met Gly Arg His Glu Thr Val Val Ser Leu Ala Ala Glu 915 920 925

Gly Gln Tyr 930

<210> 16

<211> 931

<212> PRT

<213> Caenorhabditis elegans

<400> 16

Met Arg Lys Gly Leu Arg Ala Thr Ala Ala Arg Cys Gly Leu Gly Leu 1 5 10 15

Gly Tyr Leu Leu Gln Met Leu Val Leu Pro Ala Leu Ala Leu Leu Ser 20 25 30

Ala Ser Gly Thr Gly Ser Ala Ala Gln Asp Asp Phe Phe His Glu 35 40 45

Leu Pro Glu Thr Phe Pro Ser Asp Pro Pro Glu Pro Leu Pro His Phe 50 55 60

Leu Ile Glu Pro Glu Glu Ala Tyr Ile Val Lys Asn Lys Pro Val Asn 65 70 75 80

Leu Tyr Cys Lys Ala Ser Pro Ala Thr Gln Ile Tyr Phe Lys Cys Asn 85 90 95

Ser Glu Trp Val His Gln Lys Asp His Ile Val Asp Glu Arg Val Asp 100 105 110

Glu Thr Ser Gly Leu Ile Val Arg Glu Val Ser Ile Glu Ile Ser Arg 115 120 125

Gln Gln Val Glu Glu Leu Phe Gly Pro Glu Asp Tyr Trp Cys Gln Cys 130 135 140

Val 145	Ala	Trp	Ser	Ser	Ala 150	Gly	Thr	Thr	Lys	Ser 155	Arg	Lys	Ala	Tyr	Val 160
Arg	Ile	Ala	Tyr	Leu 165	Arg	Lys	Thr	Phe	Glu 170	Gln	Glu	Pro	Leu	Gly 175	Lys
Glu	Val	Ser	Leu 180	Glu	Gln	Glu	Val	Leu 185	Leu	Gln	Cys	Arg	Pro 190	Pro	Glu
Gly	Ile	Pro 195	Val	Ala	Glu	Val	Glu 200	Trp	Leu	Lys	Asn	Glu 205	Asp	Ile	Ile
Asp	Pro 210	Val	Glu	Asp	Arg	Asn 215	Phe	Tyr	Ile	Thr	Ile 220	Asp	His	Asn	Leu
Ile 225	Ile	Lys	Gln	Ala	Arg 230	Leu	Ser	Asp	Thr	Ala 235	Asn	Tyr	Thr	Cys	Val 240
Ala	Lys	Asn	Ile	Val 245	Ala	Lys	Arg	Lys	Ser 250	Thr	Thr	Ala	Thr	Val 255	Ile
Val	Tyr	Val	Asn 260	Gly	Gly	Trp	Ser	Thr 265	Trp	Thr	Glu	Trp	Ser 270	Val	Cys
Asn	Ser	Arg 275	Cys	Gly	Arg	Gly	Tyr 280	Gln	Lys	Arg	Thr	Arg 285	Thr	Cys	Thr
Asn	Pro 290	Ala	Pro	Leu	Asn	Gly 295	Gly	Ala	Phe	Cys	Glu 300	Gly	Gln	Ser	Val
Gln 305	Lys	Ile	Ala	Cys	Thr 310	Thr	Leu	Cys	Pro	Val 315	Asp	Gly	Arg	Trp	Thr 320
Pro	Trp	Ser	Lys	Trp 325	Ser	Thr	Cys	Gly	Thr 330	Glu	Cys	Thr	His	Trp 335	Arg
Arg	Arg	Glu	Cys 340	Thr	Ala	Pro	Ala	Pro 345	Lys	Asn	Gly	Gly	Lys 350	Asp	Cys
Asp	Gly	Leu 355	Val	Leu	Gln	Ser	Lys 360	Asn	Cys	Thr	Asp	Gly 365	Leu	Cys	Met
Gln	Thr 370	Ala	Pro	Asp	Ser	Asp 375	Asp	Val	Ala	Leu	Tyr 380	Val	Gly	Ile	Val
Ile 385	Ala	Val	Ile	Val	Cys 390	Leu	Ala	Ile	Ser	Val 395	Val	Val	Ala	Leu	Phe 400
Val	Tyr	Arg	Lys	Asn 405	His	Arg	Asp	Phe	Glu 410	Ser	Asp	Ile	Ile	Asp 415	Ser
Ser	Ala	Leu	Asn 420	Gly	Gly	Phe ·	Gln	Pro 425	Val	Asn	Ile	Lys	Ala 430	Ala	Arg
Gln	Asp	Leu 435	Leu	Ala	Val	Pro	Pro 440	Asp	Leu	Thr	Ser	Ala 445	Ala	Ala	Met

Tyr Arg Gly Pro Val Tyr Ala Leu His Asp Val Ser Asp Lys Ile Pro Met Thr Asn Ser Pro Ile Leu Asp Pro Leu Pro Asn Leu Lys Ile Lys 470 475 Val Tyr Asn Thr Ser Gly Ala Val Thr Pro Gln Asp Asp Leu Ser Glu 490 Phe Thr Ser Lys Leu Ser Pro Gln Met Thr Gln Ser Leu Leu Glu Asn 500 Glu Ala Leu Ser Leu Lys Asn Gln Ser Leu Ala Arq Gln Thr Asp Pro Ser Cys Thr Ala Phe Gly Ser Phe Asn Ser Leu Gly Gly His Leu Ile 535 Val Pro Asn Ser Gly Val Ser Leu Leu Ile Pro Ala Gly Ala Ile Pro 550 555 Gln Gly Arg Val Tyr Glu Met Tyr Val Thr Val His Arg Lys Glu Thr 565 570 Met Arg Pro Pro Met Asp Asp Ser Gln Thr Leu Leu Thr Pro Val Val Ser Cys Gly Pro Pro Gly Ala Leu Leu Thr Arg Pro Val Val Leu Thr 600 Met His His Cys Ala Asp Pro Asn Thr Glu Asp Trp Lys Ile Leu Leu 615 Lys Asn Gln Ala Ala Gln Gly Gln Trp Glu Asp Val Val Val Val Gly 625 630 Glu Glu Asn Phe Thr Thr Pro Cys Tyr Ile Gln Leu Asp Ala Glu Ala 650 Cys His Ile Leu Thr Glu Asn Leu Ser Thr Tyr Ala Leu Val Gly His 660 665 Ser Thr Thr Lys Ala Ala Ala Lys Arg Leu Lys Leu Ala Ile Phe Gly 680 Pro Leu Cys Cys Ser Ser Leu Glu Tyr Ser Ile Arg Val Tyr Cys Leu 695 Asp Asp Thr Gln Asp Ala Leu Lys Glu Ile Leu His Leu Glu Arg Gln 705 710 Met Gly Gly Gln Leu Leu Glu Glu Pro Lys Ala Leu His Phe Lys Gly Ser Thr His Asn Leu Arg Leu Ser Ile His Asp Ile Ala His Ser Leu 740 745

Trp Lys Ser Lys Leu Leu Ala Lys Tyr Gln Glu Ile Pro Phe Tyr His 760 Val Trp Ser Gly Ser Gln Arg Asn Leu His Cys Thr Phe Thr Leu Glu 775 Arg Phe Ser Leu Asn Thr Val Glu Leu Val Cys Lys Leu Cys Val Arg 790 Gln Val Glu Gly Glu Gly Gln Ile Phe Gln Leu Asn Cys Thr Val Ser Glu Glu Pro Thr Gly Ile Asp Leu Pro Leu Leu Asp Pro Ala Asn Thr 825 Ile Thr Thr Val Thr Gly Pro Ser Ala Phe Ser Ile Pro Leu Pro Ile 840 Arg Gln Lys Leu Cys Ser Ser Leu Asp Ala Pro Gln Thr Arg Gly His 855 Asp Trp Arq Met Leu Ala His Lys Leu Asn Leu Asp Arq Tyr Leu Asn 870 875 Tyr Phe Ala Thr Lys Ser Ser Pro Thr Gly Val Ile Leu Asp Leu Trp 885 890 Glu Ala Gln Asn Phe Pro Asp Gly Asn Leu Ser Met Leu Ala Ala Val 905 Leu Glu Glu Met Gly Arg His Glu Thr Val Val Ser Leu Ala Ala Glu 920 Gly Gln Tyr 930 <210> 17 <211> 931 <212> PRT <213> Caenorhabditis elegans <400> 17 Met Arg Lys Gly Leu Arg Ala Thr Ala Ala Arg Cys Gly Leu Gly Leu 5 15 Gly Tyr Leu Leu Gln Met Leu Val Leu Pro Ala Leu Ala Leu Leu Ser Ala Ser Gly Thr Gly Ser Ala Ala Gln Asp Asp Asp Phe Phe His Glu Leu Pro Glu Thr Phe Pro Ser Asp Pro Pro Glu Pro Leu Pro His Phe 50 Leu Ile Glu Pro Glu Glu Ala Tyr Ile Val Lys Asn Lys Pro Val Asn 70 75

Leu Tyr Cys Lys Ala Ser Pro Ala Thr Gln Ile Tyr Phe Lys Cys Asn 90 Ser Glu Trp Val His Gln Lys Asp His Ile Val Asp Glu Arg Val Asp 100 105 Glu Thr Ser Gly Leu Ile Val Arg Glu Val Ser Ile Glu Ile Ser Arg 120 Gln Gln Val Glu Glu Leu Phe Gly Pro Glu Asp Tyr Trp Cys Gln Cys 135 Val Ala Trp Ser Ser Ala Gly Thr Thr Lys Ser Arg Lys Ala Tyr Val Arg Ile Ala Tyr Leu Arg Lys Thr Phe Glu Gln Glu Pro Leu Gly Lys 170 Glu Val Ser Leu Glu Gln Glu Val Leu Leu Gln Cys Arg Pro Pro Glu 185 Gly Ile Pro Val Ala Glu Val Glu Trp Leu Lys Asn Glu Asp Ile Ile 195 Asp Pro Val Glu Asp Arg Asn Phe Tyr Ile Thr Ile Asp His Asn Leu Ile Ile Lys Gln Ala Arg Leu Ser Asp Thr Ala Asn Tyr Thr Cys Val 225 230 235 Ala Lys Asn Ile Val Ala Lys Arg Lys Ser Thr Thr Ala Thr Val Ile Val Tyr Val Asn Gly Gly Trp Ser Thr Trp Thr Glu Trp Ser Val Cys 265 Asn Ser Arg Cys Gly Arg Gly Tyr Gln Lys Arg Thr Arg Thr Cys Thr Asn Pro Ala Pro Leu Asn Gly Gly Ala Phe Cys Glu Gly Gln Ser Val 295 Gln Lys Ile Ala Cys Thr Thr Leu Cys Pro Val Asp Gly Arg Trp Thr 305 310 315 320 Pro Trp Ser Lys Trp Ser Thr Cys Gly Thr Glu Cys Thr His Trp Arg 325 Arg Arg Glu Cys Thr Ala Pro Ala Pro Lys Asn Gly Gly Lys Asp Cys 345 Asp Gly Leu Val Leu Gln Ser Lys Asn Cys Thr Asp Gly Leu Cys Met 355 Gln Thr Ala Pro Asp Ser Asp Asp Val Ala Leu Tyr Val Gly Ile Val 370 375 380

Ile 385	Ala	Val	Ile	Val	Cys 390	Leu	Ala	Ile	Ser	Val 395	Val	Val	Ala	Leu	Phe 400
Val	Tyr	Arg	Lys	Asn 405	His	Arg	Asp	Phe	Glu 410	Ser	Asp	Ile	Ile	Asp 415	Ser
Ser	Ala	Leu	Asn 420	Gly	Gly	Phe	Gln	Pro 425	Val	Asn	Ile	Lys	Ala 430	Ala	Arg
Gln	Asp	Leu 435	Leu	Ala	Val	Pro	Pro 440	Asp	Leu	Thr	Ser	Ala 445	Ala	Ala	Met
Tyr	Arg 450	Gly	Pro	Val	Tyr	Ala 455	Leu	His	Asp	Val	Ser 460	Asp	Lys	Ile	Pro
Met 465	Thr	Asn	Ser	Pro	Ile 470	Leu	Asp	Pro	Leu	Pro 475	Asn	Leu	Lys	Ile	Lys 480
Val	Tyr	Asn	Thr	Ser 485	Gly	Ala	Val	Ser	Pro 490	Gln	Asp	Asp	Leu	Ser 495	Glu
Phe	Thr	Ser	Lys 500	Leu	Ser	Pro	Gln	Met 505	Thr	Gln	Ser	Leu	Leu 510	Glu	Asn
Glu	Ala	Leu 515	Ser	Leu	Lys	Asn	Gln 520	Ser	Leu	Ala	Arg	Gln 525	Thr	Asp	Pro
Ser	Cys 530	Thr	Ala	Phe	Gly	Ser 535	Phe	Asn	Ser	Leu	Gly 540	Gly	His	Leu	Ile
Val 545	Pro	Asn	Ser	Gly	Val 550	Ser	Leu	Leu	Ile	Pro 555	Ala	Gly	Ala	Ile	Pro 560
Gln	Gly	Arg	Val	Tyr 565	Glu	Met	Tyr	Val	Thr 570	Val	His	Arg	Lys	Glu 575	Thr
Met	Arg	Pro	Pro 580	Met	Asp	Asp	Ser	Gln 585	Thr	Leu	Leu	Thr	Pro 590	Val	Val
Ser	Cys	Gly 595	Pro	Pro	Gly	Ala	Leu 600	Leu	Thr	Arg	Pro	Val 605	Val	Leu	Thr
Met	His 610	His	Cys	Ala	Asp	Pro 615	Asn	Thr	Glu	Asp	Trp 620	Lys	Ile	Leu	Leu
Lys 625	Asn	Gln	Ala	Ala	Gln 630	Gly	Gln	Trp	Glu	Asp 635	Val	Val	Val	Val	Gly 640
Glu	Glu	Asn	Phe	Thr 645	Thr	Pro	Cys	Tyr	Ile 650	Lys	Leu	Asp	Ala	Glu 655	Ala
Cys	His	Ile	Leu 660	Thr	Glu	Asn	Leu	Ser 665	Thr	Туг	Ala	Leu	Val 670	Gly	His
Ser	Thr	Thr 675	Lys	Ala	Ala	Ala	Lys 680	Arg	Leu	Lys	Leu	Ala 685	Ile	Phe	Gly

Pro Leu Cys Cys Ser Ser Leu Glu Tyr Ser Ile Arg Val Tyr Cys Leu 690 695 700

Asp Asp Thr Gln Asp Ala Leu Lys Glu Ile Leu His Leu Glu Arg Gln 705 710 715 720

Thr Gly Gly Gln Leu Leu Glu Glu Pro Lys Ala Leu His Phe Lys Gly 725 730 735

Ser Thr His Asn Leu Arg Leu Ser Ile His Asp Ile Ala His Ser Leu
740 745 750

Trp Lys Ser Lys Leu Leu Ala Lys Tyr Gln Glu Ile Pro Phe Tyr His
755 760 765

Val Trp Ser Gly Ser Gln Arg Asn Leu His Cys Thr Phe Thr Leu Glu 770 775 780

Arg Phe Ser Leu Asn Thr Val Glu Leu Val Cys Lys Leu Cys Val Arg 785 790 795 800

Gln Val Glu Gly Glu Gly Gln Ile Phe Gln Leu Asn Cys Thr Val Ser 805 810 815

Glu Glu Pro Thr Gly Ile Asp Leu Pro Leu Leu Asp Pro Ala Asn Thr 820 825 830

Ile Thr Thr Val Thr Gly Pro Ser Ala Phe Ser Ile Pro Leu Pro Ile 835 840 845

Arg Gln Lys Leu Cys Ser Ser Leu Asp Ala Pro Gln Thr Arg Gly His 850 860

Asp Trp Arg Met Leu Ala His Lys Leu Asn Leu Asp Arg Tyr Leu Asn 865 870 875 880

Tyr Phe Ala Thr Lys Ser Ser Pro Thr Gly Val Ile Leu Asp Leu Trp 885 890 895

Glu Ala Gln Asn Phe Pro Asp Gly Asn Leu Ser Met Leu Ala Ala Val 900 905 910

Leu Glu Glu Met Gly Arg His Glu Thr Val Val Ser Leu Ala Ala Glu 915 920 925

Gly Gln Tyr 930

<210> 18

<211> 4349

<212> PRT

<213> Homo sapiens

<400> 18

Met Thr Ile Ala Leu Leu Gly Phe Ala Ile Phe Leu Leu His Cys Ala

- Thr Cys Glu Lys Pro Leu Glu Gly Ile Leu Ser Ser Ser Ala Trp His 20 25 30
- Phe Thr His Ser His Tyr Asn Ala Thr Ile Tyr Glu Asn Ser Ser Pro 35 40 45
- Lys Thr Tyr Val Glu Ser Phe Glu Lys Met Gly Ile Tyr Leu Ala Glu
  50 55 60
- Pro Gln Trp Ala Val Arg Tyr Arg Ile Ile Ser Gly Asp Val Ala Asn 65 70 75 80
- Val Phe Lys Thr Glu Glu Tyr Val Val Gly Asn Phe Cys Phe Leu Arg 85 90 95
- Ile Arg Thr Lys Ser Ser Asn Thr Ala Leu Leu Asn Arg Glu Val Arg
  100 105 110
- Asp Ser Tyr Thr Leu Ile Ile Gln Ala Thr Glu Lys Thr Leu Glu Leu 115 120 125
- Glu Ala Leu Thr Arg Val Val Val His Ile Leu Asp Gln Asn Asp Leu 130 135 140
- Met Pro Leu Lys Ser Pro Ile Cys Lys Val Thr Ala Thr Asp Ala Asp 165 170 175
- Leu Gly Gln Asn Ala Glu Phe Tyr Tyr Ala Phe Asn Thr Arg Ser Glu 180 185 190
- Met Phe Ala Ile His Pro Thr Ser Gly Val Val Thr Val Ala Gly Lys
  195 200 205
- Leu Asn Val Thr Trp Arg Gly Lys His Glu Leu Gln Val Leu Ala Val 210 215 220
- Asp Arg Met Arg Lys Ile Ser Glu Gly Asn Gly Phe Gly Ser Leu Ala 225 230 235 240
- Ala Leu Val Val His Val Glu Pro Ala Leu Arg Lys Pro Pro Ala Ile 245 250 255
- Ala Ser Val Val Val Thr Pro Pro Asp Ser Asn Asp Gly Thr Thr Tyr 260 265 270
- Ala Thr Val Leu Val Asp Ala Asn Ser Ser Gly Ala Glu Val Glu Ser 275 280 285
- Val Glu Val Val Gly Gly Asp Pro Gly Lys His Phe Lys Ala Ile Lys 290 295 300
- Ser Tyr Ala Arg Ser Asn Glu Phe Ser Leu Val Ser Val Lys Asp Ile

320 320	305	310	315	320
---------	-----	-----	-----	-----

Asn	Trp	Met	Glu	Tyr	Leu	His	Gly	Phe	Asn	Leu	Ser	Leu	Gln	Ala	Arg
				325					330					335	

- Ser Gly Ser Gly Pro Tyr Phe Tyr Ser Gln Ile Arg Gly Phe His Leu 340 345 350
- Pro Pro Ser Lys Leu Ser Ser Leu Lys Phe Glu Lys Ala Val Tyr Arg 355 360 365
- Val Gln Leu Ser Glu Phe Ser Pro Pro Gly Ser Arg Val Val Met Val 370 380
- Arg Val Thr Pro Ala Phe Pro Asn Leu Gln Tyr Val Leu Lys Pro Ser 385 390 395 400
- Ser Glu Asn Val Gly Phe Lys Leu Asn Ala Arg Thr Gly Leu Ile Thr  $405 \hspace{1.5cm} 410 \hspace{1.5cm} 415 \hspace{1.5cm}$
- Thr Thr Lys Leu Met Asp Phe His Asp Arg Ala His Tyr Gln Leu His 420 425 430
- Ile Arg Thr Ser Pro Gly Gln Ala Ser Thr Val Val Val Ile Asp Ile 435 440 445
- Val Asp Cys Asn Asn His Ala Pro Leu Phe Asn Arg Ser Ser Tyr Asp 450 455 460
- Gly Thr Leu Asp Glu Asn Ile Pro Pro Gly Thr Ser Val Leu Ala Val 465 470 475 480
- Thr Ala Thr Asp Arg Asp His Gly Glu Asn Gly Tyr Val Thr Tyr Ser 485 490 495
- Ile Ala Gly Pro Lys Ala Leu Pro Phe Ser Ile Asp Pro Tyr Leu Gly 500 505 510
- Ile Ile Ser Thr Ser Lys Pro Met Asp Tyr Glu Leu Met Lys Arg Ile 515 520 525
- Tyr Thr Phe Arg Val Arg Ala Ser Asp Trp Gly Ser Pro Phe Arg Arg 530 540
- Glu Lys Glu Val Ser Ile Phe Leu Gln Leu Arg Asn Leu Asn Asp Asn 545 550 555 560
- Gln Pro Met Phe Glu Glu Val Asn Cys Thr Gly Ser Ile Arg Gln Asp 565 570 575
- Trp Pro Val Gly Lys Ser Ile Met Thr Met Ser Ala Ile Asp Val Asp 580 585 590
- Glu Leu Gln Asn Leu Lys Tyr Glu Ile Val Ser Gly Asn Glu Leu Glu
  595 600 605
- Tyr Phe Asp Leu Asn His Phe Ser Gly Val Ile Ser Leu Lys Arg Pro

	610					615					620				
Phe 625	Ile	Asn	Leu	Thr	Ala 630	Gly	Gln	Pro	Thr	Ser 635	Tyr	Ser	Leu	Lys	Ile 640
Thr	Ala	Ser	Asp	Gly 645	Lys	Asn	Tyr	Ala	Ser 650	Pro	Thr	Thr	Leu	Asn 655	Ile
Thr	Val	Val	Lys 660	Asp	Pro	His	Phe	Glu 665	Val	Pro	Val	Thr	Cys 670	Asp	Lys
Thr	Gly	Val 675	Leu	Thr	Gln	Phe	Thr 680	Lys	Thr	Ile	Leu	His 685	Phe	Ile	Gly
Leu	Gln 690	Asn	Gln	Glu	Ser	Ser 695	Asp	Glu	Glu	Phe	Thr 700	Ser	Leu	Ser	Thr
Tyr 705	Gln	Ile	Asn	His	Tyr 710	Thr	Pro	Gln	Phe	Glu 715	Asp	His	Phe	Pro	Gln 720
Ser	Ile	Asp	Val	Leu 725	Glu	Ser	Val	Pro	Ile 730	Asn	Thr	Pro	Leu	Ala 735	Arg
Leu	Ala	Ala	Thr 740	Asp	Pro	Asp	Ala	Gly 745	Phe	Asn	Gly	Lys	Leu 750	Val	Tyr
Val	Ile	Ala 755	Asp	Gly	Asn	Glu	Glu 760	Gly	Cys	Phe	Asp	Ile 765	Glu	Leu	Glu
Thr	Gly 770	Leu	Leu	Thr	Val	Ala 775	Ala	Pro	Leu	Asp	Tyr 780	Glu	Ala	Thr	Asn
Phe 785	Tyr	Ile	Leu	Asn	Val 790	Thr	Val	Tyr	Asp	Leu 795	Gly	Thr	Pro	Gln	Lys 800
Ser	Ser	Trp	Lys	Leu 805	Leu	Thr	Val	Asn	Val 810	Lys	Asp	Trp	Asn	Asp 815	Asn
Ala	Pro	Arg	Phe 820	Pro	Pro	Gly	Gly	Tyr 825	Gln	Leu	Thr	Ile	Ser 830	Glu	Asp
Thr	Glu	Val 835	Gly	Thr	Thr	Ile	Ala 840	Glu	Leu	Thr	Thr	Lys 845	Asp	Ala	Asp
Ser	Glu 850	Asp	Asn	Gly	Arg	Val 855	Arg	Tyr	Thr	Leu	Leu 860	Ser	Pro	Thr	Glu
Lys 865	Phe	Ser	Leu	His	Pro 870	Leu	Thr	Gly	Glu	Leu 875	Val	Val	Thr	Gly	His 880
Leu	Asp	Arg	Glu	Ser 885	Glu	Pro	Arg	Tyr	Ile 890	Leu	Lys	Val	Glu	Ala 895	Arg
Asp	Gln	Pro	Ser 900	Lys	Gly	His	Gln	Leu 905	Phe	Ser	Val	Thr	Asp 910	Leu	Ile

Ile Thr Leu Glu Asp Val Asn Asp Asn Ser Pro Gln Cys Ile Thr Glu

915	920	925

- His Asn Arg Leu Lys Val Pro Glu Asp Leu Pro Pro Gly Thr Val Leu 930 935 940
- Thr Phe Leu Asp Ala Ser Asp Pro Asp Leu Gly Pro Ala Gly Glu Val 945 950 955 960
- Arg Tyr Val Leu Met Asp Gly Ala His Gly Thr Phe Arg Val Asp Leu 965 970 975
- Met Thr Gly Ala Leu Ile Leu Glu Arg Glu Leu Asp Phe Glu Arg Arg 980 985 990
- Ala Gly Tyr Asn Leu Ser Leu Trp Ala Ser Asp Gly Gly Arg Pro Leu 995 1000 1005
- Ala Arg Arg Thr Leu Cys His Val Glu Val Ile Val Leu Asp Val Asn 1010 1015 1020
- Glu Asn Leu His Pro Pro His Phe Ala Ser Phe Val His Gln Gly Gln 1025 1030 1035 1040
- Val Gln Glu Asn Ser Pro Ser Gly Thr Gln Val Ile Val Val Ala Ala
  1045 1050 1055
- Gln Asp Asp Ser Gly Leu Asp Gly Glu Leu Gln Tyr Phe Leu Arg 1060 1065 1070
- Ala Gly Thr Gly Leu Ala Ala Phe Ser Ile Asn Gln Asp Thr Gly Met
  1075 1080 1085
- Ile Gln Thr Leu Ala Pro Leu Asp Arg Glu Phe Ala Ser Tyr Tyr Trp 1090 1095 1100
- Leu Thr Val Leu Ala Val Asp Arg Gly Ser Val Pro Leu Ser Ser Val 1105 1110 1115 1120
- Thr Glu Val Tyr Ile Glu Val Thr Asp Ala Asn Asp Asn Pro Pro Gln
  1125 1130 1135
- Met Ser Gln Ala Val Phe Tyr Pro Ser Ile Gln Glu Asp Ala Pro Val 1140 1145 1150
- Gly Thr Ser Val Leu Gln Leu Asp Ala Trp Asp Pro Asp Ser Ser Ser 1155 1160 1165
- Lys Gly Lys Leu Thr Phe Asn Ile Thr Ser Gly Asn Tyr Met Gly Phe 1170 1175 1180
- Phe Met Ile His Pro Val Thr Gly Leu Leu Ser Thr Ala Gln Gln Leu 1185 1190 1195 1200
- Asp Arg Glu Asn Lys Asp Glu His Ile Leu Glu Val Thr Val Leu Asp 1205 1210 1215
- Asn Gly Glu Pro Ser Leu Lys Ser Thr Ser Arg Val Val Gly Ile

- Leu Asp Val Asn Asp Asn Pro Pro Ile Phe Ser His Lys Leu Phe Asn 1235 1240 1245
- Val Arg Leu Pro Glu Arg Leu Ser Pro Val Ser Pro Gly Pro Val Tyr 1250 1260
- Arg Leu Val Ala Ser Asp Leu Asp Glu Gly Leu Asn Gly Arg Val Thr 1265 1270 1275 1280
- Tyr Ser Ile Glu Asp Ser Asp Glu Glu Ala Phe Ser Ile Asp Leu Val 1285 1290 1295
- Thr Gly Val Val Ser Ser Ser Ser Thr Phe Thr Ala Gly Glu Tyr Asn 1300 1305 1310
- Ile Leu Thr Ile Lys Ala Thr Asp Ser Gly Gln Pro Pro Leu Ser Ala 1315 1320 1325
- Ser Val Arg Leu His Ile Glu Trp Ile Pro Trp Pro Arg Pro Ser Ser 1330 1335 1340
- Ile Pro Leu Ala Phe Asp Glu Thr Tyr Tyr Ser Phe Thr Val Met Glu 1345 1350 1355 1360
- Thr Asp Pro Val Asn His Met Val Gly Val Ile Ser Val Glu Gly Arg 1365 1370 1375
- Pro Gly Leu Phe Trp Phe Asn Ile Ser Gly Gly Asp Lys Asp Met Asp 1380 1385 1390
- Phe Asp Ile Glu Lys Thr Thr Gly Ser Ile Val Ile Ala Arg Pro Leu 1395 1400 1405
- Asp Thr Arg Arg Arg Ser Asn Tyr Asn Leu Thr Val Glu Val Thr Asp 1410 1415 1420
- Gly Ser Arg Thr Ile Ala Thr Gln Val His Ile Phe Met Ile Ala Asn 1425 1430 1435 1440
- Ile Asn His His Arg Pro Gln Phe Leu Glu Thr Arg Tyr Glu Val Arg 1445 1450 1455
- Val Pro Gln Asp Thr Val Pro Gly Val Glu Leu Leu Arg Val Gln Ala 1460 1465 1470
- Ile Asp Gln Asp Lys Gly Lys Ser Leu Ile Tyr Thr Ile His Gly Ser 1475 1480 1485
- Gln Asp Pro Gly Ser Ala Ser Leu Phe Gln Leu Asp Pro Ser Ser Gly 1490 1495 1500
- Val Leu Val Thr Val Gly Lys Leu Asp Leu Gly Ser Gly Pro Ser Gln 1505 1510 1515 1520
- His Thr Leu Thr Val Met Val Arg Asp Gln Glu Ile Pro Ile Lys Arg

- Asn Phe Val Trp Val Thr Ile His Val Glu Asp Gly Asn Leu His Pro  $1540 \hspace{1cm} 1545 \hspace{1cm} 1550$
- Pro Arg Phe Thr Gln Leu His Tyr Glu Ala Ser Val Pro Asp Thr Ile 1555 1560 1565
- Ala Pro Gly Thr Glu Leu Leu Gln Val Arg Ala Met Asp Ala Asp Arg 1570 1575 1580
- Gly Val Asn Ala Glu Val His Tyr Ser Leu Leu Lys Gly Asn Ser Glu 1585 1590 1595 1600
- Gly Phe Phe Asn Ile Asn Ala Leu Leu Gly Ile Ile Thr Leu Ala Gln 1605 1610 1615
- Lys Leu Asp Gln Ala Asn His Ala Pro His Thr Leu Thr Val Lys Ala 1620 1625 1630
- Glu Asp Gln Gly Ser Pro Gln Trp His Asp Leu Ala Thr Val Ile Ile 1635 1640 1645
- His Val Tyr Pro Ser Asp Arg Ser Ala Pro Ile Phe Ser Lys Ser Glu 1650 1655 1660
- Tyr Phe Val Glu Ile Pro Glu Ser Ile Pro Val Gly Ser Pro Ile Leu 1665 1670 1675 1680
- Leu Val Ser Ala Met Ser Pro Ser Glu Val Thr Tyr Glu Leu Arg Glu 1685 1690 1695
- Gly Asn Lys Asp Gly Val Phe Ser Met Asn Ser Tyr Ser Gly Leu Ile 1700 1705 1710
- Ser Thr Gln Lys Lys Leu Asp His Glu Lys Ile Ser Ser Tyr Gln Leu 1715 1720 1725
- Lys Ile Arg Gly Ser Asn Met Ala Gly Ala Phe Thr Asp Val Met Val 1730 1735 1740
- Val Val Asp Ile Ile Asp Glu Asn Asp Asn Ala Pro Met Phe Leu Lys 1745 1750 1755 1760
- Ser Thr Phe Val Gly Gln Ile Ser Glu Ala Ala Pro Leu Tyr Ser Met 1765 1770 1775
- Ile Met Asp Lys Asn Asn Pro Phe Val Ile His Ala Ser Asp Ser 1780 1785 1790
- Asp Lys Glu Ala Asn Ser Leu Leu Val Tyr Lys Ile Leu Glu Pro Glu 1795 1800 1805
- Ala Leu Lys Phe Phe Lys Ile Asp Pro Ser Met Gly Thr Leu Thr Ile 1810 1815 1820
- Val Ser Glu Met Asp Tyr Glu Ser Met Pro Ser Phe Gln Phe Cys Val

- Tyr Val His Asp Gln Gly Ser Pro Val Leu Phe Ala Pro Arg Pro Ala 1845 1850 1855
- Gln Val Ile Ile His Val Arg Asp Val Asn Asp Ser Pro Pro Arg Phe 1860 1865 1870
- Ser Glu Gln Ile Tyr Glu Val Ala Ile Val Gly Pro Ile His Pro Gly 1875 1880 1885
- Met Glu Leu Met Val Arg Ala Ser Asp Glu Asp Ser Glu Val Asn 1890 1895 1900
- Tyr Ser Ile Lys Thr Gly Asn Ala Asp Glu Ala Val Thr Ile His Pro 1905 1910 1915 1920
- Val Thr Gly Ser Ile Ser Val Leu Asn Pro Ala Phe Leu Gly Leu Ser 1925 1930 1935
- Arg Lys Leu Thr Ile Arg Ala Ser Asp Gly Leu Tyr Gln Asp Thr Ala 1940 1945 1950
- Leu Val Lys Ile Ser Leu Thr Gln Val Leu Asp Lys Ser Leu Gln Phe 1955 1960 1965
- Asp Gln Asp Val Tyr Trp Ala Ala Val Lys Glu Asn Leu Gln Asp Arg 1970 1975 1980
- Lys Ala Leu Val Ile Leu Gly Ala Gln Gly Asn His Leu Asn Asp Thr 1985 1990 1995 2000
- Leu Ser Tyr Phe Leu Leu Asn Gly Thr Asp Met Phe His Met Val Gln 2005 2010 2015
- Ser Ala Gly Val Leu Gln Thr Arg Gly Val Ala Phe Asp Arg Glu Gln 2020 2025 2030
- Gln Asp Thr His Glu Leu Ala Val Glu Val Arg Asp Asn Arg Thr Pro 2035 2040 2045
- Gln Arg Val Ala Gln Gly Leu Val Arg Val Ser Ile Glu Asp Val Asn 2050 2055 2060
- Asp Asn Pro Pro Lys Phe Lys His Leu Pro Tyr Tyr Thr Ile Ile Gln 2065 2070 2075 2080
- Asp Gly Thr Glu Pro Gly Asp Val Leu Phe Gln Val Ser Ala Thr Asp 2085 2090 2095
- Glu Asp Leu Gly Thr Asn Gly Ala Val Thr Tyr Glu Phe Ala Glu Asp 2100 2105 2110
- Tyr Thr Tyr Phe Arg Ile Asp Pro Tyr Leu Gly Asp Ile Ser Leu Lys 2115 2120 2125
- Lys Pro Phe Asp Tyr Gln Ala Leu Asn Lys Tyr His Leu Lys Val Ile

- Ala Arg Asp Gly Gly Thr Pro Ser Leu Gln Ser Glu Glu Glu Val Leu 2145 2150 2155 2160
- Val Thr Val Arg Asn Lys Ser Asn Pro Leu Phe Gln Ser Pro Tyr Tyr 2165 2170 2175
- Lys Val Arg Val Pro Glu Asn Ile Thr Leu Tyr Thr Pro Ile Leu His 2180 2185 2190
- Thr Gln Ala Arg Ser Pro Glu Gly Leu Arg Leu Ile Tyr Asn Ile Val 2195 2200 2205
- Glu Glu Glu Pro Leu Met Leu Phe Thr Thr Asp Phe Lys Thr Gly Val 2210 2215 2220
- Leu Thr Val Thr Gly Pro Leu Asp Tyr Glu Ser Lys Thr Lys His Val 2225 2230 2235 2240
- Phe Thr Val Arg Ala Thr Asp Thr Ala Leu Gly Ser Phe Ser Glu Ala 2245 2250 2255
- Thr Val Glu Val Leu Val Glu Asp Val Asn Asp Asn Pro Pro Thr Phe 2260 2265 2270
- Ser Gln Leu Val Tyr Thr Thr Ser Ile Ser Glu Gly Leu Pro Ala Gln 2275 2280 2285
- Thr Pro Val Ile Gln Leu Leu Ala Ser Asp Gln Asp Ser Gly Arg Asn 2290 2295 2300
- Arg Asp Val Ser Tyr Gln Ile Val Glu Asp Gly Ser Asp Val Ser Lys 2305 2310 2315 2320
- Phe Phe Gln Ile Asn Gly Ser Thr Gly Glu Met Ser Thr Val Gln Glu 2325 2330 2335
- Leu Asp Tyr Glu Ala Gln Gln His Phe His Val Lys Val Arg Ala Met 2340 2345 2350
- Asp Lys Gly Asp Pro Pro Leu Thr Gly Glu Thr Leu Val Val Val Asn 2355 2360 2365
- Val Ser Asp Ile Asn Asp Asn Pro Pro Glu Phe Arg Gln Pro Gln Tyr 2370 2375 2380
- Glu Ala Asn Val Ser Glu Leu Ala Thr Cys Gly His Leu Val Leu Lys 2385 2390 2395 2400
- Val Gln Ala Ile Asp Pro Asp Ser Arg Asp Thr Ser Arg Leu Glu Tyr 2405 2410 2415
- Leu Ile Leu Ser Gly Asn Gln Asp Arg His Phe Phe Ile Asn Ser Ser 2420 2425 2430
- Ser Gly Ile Ile Ser Met Phe Asn Leu Cys Lys Lys His Leu Asp Ser

2435 2440 2445

- Ser Tyr Asn Leu Arg Val Gly Ala Ser Asp Gly Val Phe Arg Ala Thr 2450 2455 2460
- Val Pro Val Tyr Ile Asn Thr Thr Asn Ala Asn Lys Tyr Ser Pro Glu 2465 2470 2475 2480
- Phe Gln Gln His Leu Tyr Glu Ala Glu Leu Ala Glu Asn Ala Met Val 2485 2490 2495
- Gly Thr Lys Val Ile Asp Leu Leu Ala Ile Asp Lys Asp Ser Gly Pro 2500 2505 2510
- Tyr Gly Thr Ile Asp Tyr Thr Ile Ile Asn Lys Leu Ala Ser Glu Lys 2515 2520 2525
- Phe Ser Ile Asn Pro Asn Gly Gln Ile Ala Thr Leu Gln Lys Leu Asp 2530 2540
- Arg Glu Asn Ser Thr Glu Arg Val Ile Ala Ile Lys Val Met Ala Arg 2545 2550 2560
- Asp Gly Gly Gly Arg Val Ala Phe Cys Thr Val Lys Ile Ile Leu Thr 2565 2570 2575
- Asp Glu Asn Asp Asn Pro Pro Gln Phe Lys Ala Ser Glu Tyr Thr Val 2580 2585 2590
- Ser Ile Gln Ser Asn Val Ser Lys Asp Ser Pro Val Ile Gln Val Leu 2595 2600 2605
- Ala Tyr Asp Ala Asp Glu Gly Gln Asn Ala Asp Val Thr Tyr Ser Val 2610 2615 2620
- Asn Pro Glu Asp Leu Val Lys Asp Val Ile Glu Ile Asn Pro Val Thr 2625 2630 2635 2640
- Gly Val Val Lys Val Lys Asp Ser Leu Val Gly Leu Glu Asn Gln Thr 2645 2650 2655
- Leu Asp Phe Phe Ile Lys Ala Gln Asp Gly Gly Pro Pro His Trp Asn 2660 2665 2670
- Ser Leu Val Pro Val Arg Leu Gln Val Val Pro Lys Lys Val Ser Leu 2675 2680 2685
- Pro Lys Phe Ser Glu Pro Leu Tyr Thr Phe Ser Ala Pro Glu Asp Leu 2690 2695 2700
- Pro Glu Gly Ser Glu Ile Gly Ile Val Lys Ala Val Ala Ala Gln Asp 2705 2710 2715 2720
- Pro Val Ile Tyr Ser Leu Val Arg Gly Thr Thr Pro Glu Ser Asn Lys 2725 2730 2735
- Asp Gly Val Phe Ser Leu Asp Pro Asp Thr Gly Val Ile Lys Val Arg

- Lys Pro Met Asp His Glu Ser Thr Lys Leu Tyr Gln Ile Asp Val Met
- Ala His Cys Leu Gln Asn Thr Asp Val Val Ser Leu Val Ser Val Asn
- Ile Gln Val Gly Asp Val Asn Asp Asn Arg Pro Val Phe Glu Ala Asp
- Pro Tyr Lys Ala Val Leu Thr Glu Asn Met Pro Val Gly Thr Ser Val
- Ile Gln Val Thr Ala Ile Asp Lys Asp Thr Gly Arg Asp Gly Gln Val
- Ser Tyr Arg Leu Ser Ala Asp Pro Gly Ser Asn Val His Glu Leu Phe
- Ala Ile Asp Ser Glu Ser Gly Trp Ile Thr Thr Leu Gln Glu Leu Asp
- Cys Glu Thr Cys Gln Thr Tyr His Phe His Val Val Ala Tyr Asp His
- Gly Gln Thr Ile Gln Leu Ser Ser Gln Ala Leu Val Gln Val Ser Ile
- Thr Asp Glu Asn Asp Asn Ala Pro Arg Phe Ala Ser Glu Glu Tyr Arg
- Gly Ser Val Val Glu Asn Ser Glu Pro Gly Glu Leu Val Ala Thr Leu
- Lys Thr Leu Asp Ala Asp Ile Ser Glu Gln Asn Arg Gln Val Thr Cys
- Tyr Ile Thr Glu Gly Asp Pro Leu Gly Gln Phe Gly Ile Ser Gln Val
- Gly Asp Glu Trp Arg Ile Ser Ser Arg Lys Thr Leu Asp Arg Glu His
- Thr Ala Lys Tyr Leu Leu Arg Val Thr Ala Ser Asp Gly Lys Phe Gln
- Ala Ser Val Thr Val Glu Ile Phe Val Leu Asp Val Asn Asp Asn Ser
- Pro Gln Cys Ser Gln Leu Leu Tyr Thr Gly Lys Val His Glu Asp Val
- Phe Pro Gly His Phe Ile Leu Lys Val Ser Ala Thr Asp Leu Asp Thr
- Asp Thr Asn Ala Gln Ile Thr Tyr Ser Leu His Gly Pro Gly Ala His

- Glu Phe Lys Leu Asp Pro His Thr Gly Glu Leu Thr Thr Leu Thr Ala 3060 3065 3070
- Leu Asp Arg Glu Arg Lys Asp Val Phe Asn Leu Val Ala Lys Ala Thr 3075 3080 3085
- Asp Gly Gly Arg Ser Cys Gln Ala Asp Ile Thr Leu His Val Glu 3090 3095 3100
- Asp Val Asn Asp Asn Ala Pro Arg Phe Phe Pro Ser His Cys Ala Val 3105 3110 3120
- Ala Val Phe Asp Asn Thr Thr Val Lys Thr Pro Val Ala Val Val Phe 3125 3130 3135
- Ala Arg Asp Pro Asp Gln Gly Ala Asn Ala Gln Val Val Tyr Ser Leu 3140 3145 3150
- Pro Asp Ser Ala Glu Gly His Phe Ser Ile Asp Ala Thr Thr Gly Val 3155 3160 3165
- Ile Arg Leu Glu Lys Pro Leu Gln Val Arg Pro Gln Ala Pro Leu Glu 3170 3175 3180
- Leu Thr Val Arg Ala Ser Asp Leu Gly Thr Pro Ile Pro Leu Ser Thr 3185 3190 3195 3200
- Leu Gly Thr Val Thr Val Ser Val Val Gly Leu Glu Asp Tyr Leu Pro 3205 3210 3215
- Val Phe Leu Asn Thr Glu His Ser Val Gln Val Pro Glu Asp Ala Pro 3220 3225 3230
- Pro Gly Thr Glu Val Leu Gln Leu Ala Thr Leu Thr Arg Pro Gly Ala 3235 3240 3245
- Glu Lys Thr Gly Tyr Arg Val Val Ser Gly Asn Glu Gln Gly Arg Phe 3250 3255 3260
- Arg Leu Asp Ala Arg Thr Gly Ile Leu Tyr Val Asn Ala Ser Leu Asp 3265 3270 3275 3280
- Phe Glu Thr Ser Pro Lys Tyr Phe Leu Ser Ile Glu Cys Ser Arg Lys 3285 3290 3295
- Ser Ser Ser Ser Leu Ser Asp Val Thr Thr Val Met Val Asn Ile Thr 3300 3305 3310
- Asp Val Asn Glu His Arg Pro Gln Phe Pro Gln Asp Pro Tyr Ser Thr 3315 3320 3325
- Arg Val Leu Glu Asn Ala Leu Val Gly Asp Val Ile Leu Thr Val Ser 3330 3335 3340
- Ala Thr Asp Glu Asp Gly Pro Leu Asn Ser Asp Ile Thr Tyr Ser Leu

3345 3350 3355 3360

Ile Gly Gly Asn Gln Leu Gly His Phe Thr Ile His Pro Lys Lys Gly 3365 3370 3375

Glu Leu Gln Val Ala Lys Ala Leu Asp Arg Glu Gln Ala Ser Ser Tyr 3380 3385 3390

Ser Leu Lys Leu Arg Ala Thr Asp Ser Gly Gln Pro Pro Leu His Glu 3395 3400 3405

Asp Thr Asp Ile Ala Ile Gln Val Ala Asp Val Asn Asp Asn Pro Pro 3410 3415 3420

Arg Phe Phe Gln Leu Asn Tyr Ser Thr Thr Val Gln Glu Asn Ser Pro 3425 3430 3435 3440

Ile Gly Ser Lys Val Leu Gln Leu Ile Leu Ser Asp Pro Asp Ser Pro 3445 3450 3455

Glu Asn Gly Pro Pro Tyr Ser Phe Arg Ile Thr Lys Gly Asn Asn Gly 3460 3465 3470

Ser Ala Phe Arg Val Thr Pro Asp Gly Trp Leu Val Thr Ala Glu Gly 3475 3480 3485

Leu Ser Arg Arg Ala Gln Glu Trp Tyr Gln Leu Gln Ile Gln Ala Ser 3490 3495 3500

Asp Ser Gly Ile Pro Pro Leu Ser Ser Leu Thr Ser Val Arg Val His 3505 3510 3515 3520

Val Thr Glu Gln Ser His Tyr Ala Pro Ser Ala Leu Pro Leu Glu Ile 3525 3530 3535

Phe Ile Thr Val Gly Glu Asp Glu Phe Gln Gly Gly Met Val Gly Lys 3540 3545 3550

Ile His Ala Thr Asp Arg Asp Pro Gln Asp Thr Leu Thr Tyr Ser Leu 3555 3560 3565

Ala Glu Glu Glu Thr Leu Gly Arg His Phe Ser Val Gly Ala Pro Asp 3570 3575 3580

Gly Lys Ile Ile Ala Ala Gln Gly Leu Pro Arg Gly His Tyr Ser Phe 3585 3590 3595 3600

Asn Val Thr Val Ser Asp Gly Thr Phe Thr Thr Thr Ala Gly Val His 3605 3610 3615

Val Tyr Val Trp His Val Gly Gln Glu Ala Leu Gln Gln Ala Met Trp 3620 3625 3630

Met Gly Phe Tyr Gln Leu Thr Pro Glu Glu Leu Val Ser Asp His Trp 3635 3640 3645

Arg Asn Leu Gln Arg Phe Leu Ser His Lys Leu Asp Ile Lys Arg Ala

- Asn Ile His Leu Ala Ser Leu Gln Pro Ala Glu Ala Val Ala Gly Val 3665 3670 3680
- Asp Val Leu Leu Val Phe Glu Gly His Ser Gly Thr Phe Tyr Glu Phe 3685 3690 3695
- Gln Glu Leu Ala Ser Ile Ile Thr His Ser Ala Lys Glu Met Glu His 3700 3705 3710
- Ser Val Gly Val Gln.Met Arg Ser Ala Met Pro Met Val Pro Cys Gln 3715 3720 3725
- Gly Pro Thr Cys Gln Gly Gln Ile Cys His Asn Thr Val His Leu Asp 3730 3740
- Pro Lys Val Gly Pro Thr Tyr Ser Thr Ala Arg Leu Ser Ile Leu Thr 3745 3750 3755 3760
- Pro Arg His His Leu Gln Arg Ser Cys Ser Cys Asn Gly Thr Ala Thr 3765 3770 3775
- Arg Phe Ser Gly Gln Ser Tyr Val Arg Tyr Arg Ala Pro Ala Ala Arg 3780 3785 3790
- Asn Trp His Ile His Phe Tyr Leu Lys Thr Leu Gln Pro Gln Ala Ile 3795 3800 3805
- Leu Leu Phe Thr Asn Glu Thr Ala Ser Val Ser Leu Lys Leu Ala Ser 3810 3815 3820
- Gly Val Pro Gln Leu Glu Tyr His Cys Leu Gly Gly Phe Tyr Gly Asn 3825 3830 3835 3840
- Leu Ser Ser Gln Arg His Val Asn Asp His Glu Trp His Ser Ile Leu 3845 3850 3855
- Val Glu Met Asp Ala Ser Ile Arg Leu Met Val Asp Ser Met Gly 3860 3865 3870
- Asn Thr Ser Leu Val Val Pro Glu Asn Cys Arg Gly Leu Arg Pro Glu 3875 3880 3885
- Arg His Leu Leu Cly Gly Leu Ile Leu Leu His Ser Ser Asn 3890 3895 3900
- Val Ser Gln Gly Phe Glu Gly Cys Leu Asp Ala Val Val Val Asn Glu 3905 3910 3915 3920
- Glu Ala Leu Asp Leu Leu Ala Pro Gly Lys Thr Val Ala Gly Leu Leu 3925 3930 3935
- Glu Thr Gln Ala Leu Thr Gln Cys Cys Leu His Ser Asp Tyr Cys Ser 3940 3945 3950
- Gln Asn Thr Cys Leu Asn Gly Gly Lys Cys Ser Trp Thr His Gly Ala

- Gly Tyr Val Cys Lys Cys Pro Pro Gln Phe Ser Gly Lys His Cys Glu 3970 3975 3980
- Gln Gly Arg Glu Asn Cys Thr Phe Ala Pro Cys Leu Glu Gly Gly Thr 3985 3990 3995 4000
- Cys Ile Leu Ser Pro Lys Gly Ala Ser Cys Asn Cys Pro His Pro Tyr 4005 4010 4015
- Thr Gly Asp Arg Cys Glu Met Glu Ala Arg Gly Cys Ser Glu Gly His
  4020 4025 4030
- Cys Leu Val Thr Pro Glu Ile Gln Arg Gly Asp Trp Gly Gln Glu 4035 4040 4045
- Leu Leu Ile Ile Thr Val Ala Val Ala Phe Ile Ile Ile Ser Thr Val 4050 4055 4060
- Gly Leu Leu Phe Tyr Cys Arg Arg Cys Lys Ser His Lys Pro Val Ala 4065 4070 4075 4080
- Met Glu Asp Pro Asp Leu Leu Ala Arg Ser Val Gly Val Asp Thr Gln
  4085 4090 4095
- Ala Met Pro Ala Ile Glu Leu Asn Pro Leu Ser Ala Ser Ser Cys Asn 4100 4105 4110
- Asn Leu Asn Gln Pro Glu Pro Ser Lys Ala Ser Val Pro Asn Glu Leu 4115 4120 4125
- Val Thr Phe Gly Pro Asn Ser Lys Gln Arg Pro Val Val Cys Ser Val 4130 4135 4140
- Pro Pro Arg Leu Pro Pro Ala Ala Val Pro Ser His Ser Asp Asn Glu 4145 4150 4155 4160
- Pro Val Ile Lys Arg Thr Trp Ser Ser Glu Glu Met Val Tyr Pro Gly 4165 4170 4175
- Gly Ala Met Val Trp Pro Pro Thr Tyr Ser Arg Asn Glu Arg Trp Glu 4180 4185 4190
- Tyr Pro His Ser Glu Val Thr Gln Gly Pro Leu Pro Pro Ser Ala His 4195 4200 4205
- Arg His Ser Thr Pro Val Val Met Pro Glu Pro Asn Gly Leu Tyr Gly 4210 4215 4220
- Gly Phe Pro Phe Pro Leu Glu Met Glu Asn Lys Arg Ala Pro Leu Pro 4225 4230 4235 4240
- Pro Arg Tyr Ser Asn Gln Asn Leu Glu Asp Leu Met Pro Ser Arg Pro 4245 4250 4255
- Pro Ser Pro Arg Glu Arg Leu Val Ala Pro Cys Leu Asn Glu Tyr Thr

Ala Ile Ser Tyr Tyr His Ser Gln Phe Arg Gln Gly Gly Gly Pro 4275 4280 4285

Cys Leu Ala Asp Gly Gly Tyr Lys Gly Val Gly Met Arg Leu Ser Arg 4290 4295 4300

Ala Gly Pro Ser Tyr Ala Val Cys Glu Val Glu Gly Ala Pro Leu Ala 4305 4310 4315 4320

Gly Gln Gly Gln Pro Arg Val Pro Pro Asn Tyr Glu Gly Ser Asp Met 4325 4330 4335

Val Glu Ser Asp Tyr Gly Ser Cys Glu Glu Val Met Phe 4340 4345

<210> 19

<211> 4349

<212> PRT

<213> Homo sapiens

<400> 19

Met Thr Ile Ala Leu Leu Gly Phe Ala Ile Phe Leu Leu His Cys Ala 1 5 10 15

Thr Cys Glu Lys Pro Leu Glu Gly Ile Leu Ser Ser Ser Ala Trp His 20 25 30

Phe Thr His Ser His Tyr Asn Ala Thr Ile Tyr Glu Asn Ser Ser Pro 35 40 45

Lys Thr Tyr Val Glu Ser Phe Glu Lys Met Gly Ile Tyr Leu Ala Glu 50 55 60

Pro Gln Trp Ala Val Arg Tyr Arg Ile Ile Ser Gly Asp Val Ala Asn 65 70 75 80

Val Phe Lys Thr Glu Glu Tyr Val Val Gly Asn Phe Cys Phe Leu Arg 85 90 95

Ile Arg Thr Lys Ser Ser Asn Thr Ala Leu Leu Asn Arg Glu Val Arg
100 105 110

Asp Ser Tyr Thr Leu Ile Ile Gln Ala Thr Glu Lys Thr Leu Glu Leu 115 120 125

Glu Ala Leu Thr Arg Val Val Val His Ile Leu Asp Gln Asn Asp Leu 130 135 140

Lys Pro Leu Phe Ser Pro Pro Ser Tyr Arg Val Thr Ile Ser Glu Asp 145 150 155 160

Met Pro Leu Lys Ser Pro Ile Cys Lys Val Thr Ala Thr Asp Ala Asp 165 170 175

Leu Gly Gln Asn Ala Glu Phe Tyr Tyr Ala Phe Asn Thr Arg Ser Glu Met Phe Ala Ile His Pro Thr Ser Gly Val Val Thr Val Ala Gly Lys 200 Leu Asn Val Thr Trp Arg Gly Lys His Glu Leu Gln Val Leu Ala Val 215 Asp Arg Met Arg Lys Ile Ser Glu Gly Asn Gly Phe Gly Ser Leu Ala Ala Leu Val Val His Val Glu Pro Ala Leu Arg Lys Pro Pro Ala Ile 245 250 Ala Ser Val Val Thr Pro Pro Asp Ser Asn Asp Gly Thr Thr Tyr 265 Ala Thr Val Leu Val Asp Ala Asn Ser Ser Gly Ala Glu Val Glu Ser 275 280 Val Glu Val Val Gly Gly Asp Pro Gly Lys His Phe Lys Ala Ile Lys 295 Ser Tyr Ala Arg Ser Asn Glu Phe Ser Leu Val Ser Val Lys Asp Ile Asn Trp Met Glu Tyr Leu His Gly Phe Asn Leu Ser Leu Gln Ala Arg 325 330 Ser Gly Ser Gly Pro Tyr Phe Tyr Ser Gln Ile Arg Gly Phe His Leu 345 Pro Pro Ser Lys Leu Ser Ser Leu Lys Phe Glu Lys Ala Val Tyr Arg Val Gln Leu Ser Glu Phe Ser Pro Pro Gly Ser Arg Val Val Met Val 375 Arg Val Thr Pro Ala Phe Pro Asn Leu Gln Tyr Val Leu Lys Pro Ser 385 390 Ser Glu Asn Val Gly Phe Lys Leu Asn Ala Arg Thr Gly Leu Ile Thr Thr Thr Lys Leu Met Asp Phe His Asp Arg Ala His Tyr Gln Leu His 425 Ile Arg Thr Ser Pro Gly Gln Ala Ser Thr Val Val Ile Asp Ile Val Asp Cys Asn Asn His Ala Pro Leu Phe Asn Arg Ser Ser Tyr Asp 455 Gly Thr Leu Asp Glu Asn Ile Pro Pro Gly Thr Ser Val Leu Ala Val 465 470 475 480

Thr Ala Thr Asp Arg Asp His Gly Glu Asn Gly Tyr Val Thr Tyr Ser Ile Ala Gly Pro Lys Ala Leu Pro Phe Ser Ile Asp Pro Tyr Leu Gly 500 505 Ile Ile Ser Thr Ser Lys Pro Met Asp Tyr Glu Leu Met Lys Arg Ile 520 Tyr Thr Phe Arg Val Arg Ala Ser Asp Trp Gly Ser Pro Phe Arg Arg 530 535 Glu Lys Glu Val Ser Ile Phe Leu Gln Leu Arg Asn Leu Asn Asp Asn Gln Pro Met Phe Glu Glu Val Asn Cys Thr Gly Ser Ile Arq Gln Asp 570 Trp Pro Val Gly Lys Ser Ile Met Thr Met Ser Ala Ile Asp Val Asp 580 585 Glu Leu Gln Asn Leu Lys Tyr Glu Ile Val Ser Gly Asn Glu Leu Glu 600 Tyr Phe Asp Leu Asn His Phe Ser Gly Val Ile Ser Leu Lys Arg Pro 610 Phe Ile Asn Leu Thr Ala Gly Gln Pro Thr Ser Tyr Ser Leu Lys Ile Thr Ala Ser Asp Gly Lys Asn Tyr Ala Ser Pro Thr Thr Leu Asn Ile 650 Thr Val Val Lys Asp Pro His Phe Glu Val Pro Val Thr Cys Asp Lys 660 665 Thr Gly Val Leu Thr Gln Phe Thr Lys Thr Ile Leu His Phe Ile Gly 680 Leu Gln Asn Gln Glu Ser Ser Asp Glu Glu Phe Thr Ser Leu Ser Thr 690 Tyr Gln Ile Asn His Tyr Thr Pro Gln Phe Glu Asp His Phe Pro Gln 710 Ser Ile Asp Val Leu Glu Ser Val Pro Ile Asn Thr Pro Leu Ala Arg 725 730 Leu Ala Ala Thr Asp Pro Asp Ala Gly Phe Asn Gly Lys Leu Val Tyr 740 Val Ile Ala Asp Gly Asn Glu Glu Gly Cys Phe Asp Ile Glu Leu Glu Thr Gly Leu Leu Thr Val Ala Ala Pro Leu Asp Tyr Glu Ala Thr Asn 770 775

Phe Tyr Ile Leu Asn Val Thr Val Tyr Asp Leu Gly Thr Pro Gln Lys 790 Ser Ser Trp Lys Leu Leu Thr Val Asn Val Lys Asp Trp Asn Asp Asn 805 810 Ala Pro Arg Phe Pro Pro Gly Gly Tyr Gln Leu Thr Ile Ser Glu Asp 825 Thr Glu Val Gly Thr Thr Ile Ala Glu Leu Thr Thr Lys Asp Ala Asp 835 840 845 Ser Glu Asp Asn Gly Arg Val Arg Tyr Thr Leu Leu Ser Pro Thr Glu Lys Phe Ser Leu His Pro Leu Thr Gly Glu Leu Val Val Thr Gly His 875 Leu Asp Arg Glu Ser Glu Pro Arg Tyr Ile Leu Lys Val Glu Ala Arg 885 890 Asp Gln Pro Ser Lys Gly His Gln Leu Phe Ser Val Thr Asp Leu Ile 905 Ile Thr Leu Glu Asp Val Asn Asp Asn Ser Pro Gln Cys Ile Thr Glu 915 920 His Asn Arg Leu Lys Val Pro Glu Asp Leu Pro Pro Gly Thr Val Leu Thr Phe Leu Asp Ala Ser Asp Pro Asp Leu Gly Pro Ala Gly Glu Val 950 955 Arg Tyr Val Leu Met Asp Gly Ala His Gly Thr Phe Arg Val Asp Leu 970 965 Met Thr Gly Ala Leu Ile Leu Glu Arg Glu Leu Asp Phe Glu Arg Arg 985 Ala Gly Tyr Asn Leu Ser Leu Trp Ala Ser Asp Gly Gly Arg Pro Leu Ala Arg Arg Thr Leu Cys His Val Glu Val Ile Val Leu Asp Val Asn 1015 1020 Glu Asn Leu His Pro Pro His Phe Ala Ser Phe Val His Gln Gly Gln 1030 1035 Val Gln Glu Asn Ser Pro Ser Gly Thr Gln Val Ile Val Val Ala Ala 1045 1050 1055 Gln Asp Asp Ser Gly Leu Asp Gly Glu Leu Gln Tyr Phe Leu Arg 1065

Ala Gly Thr Gly Leu Ala Ala Phe Ser Ile Asn Gln Asp Thr Gly Met

- Ile Gln Thr Leu Ala Pro Leu Asp Arg Glu Phe Ala Ser Tyr Tyr Trp 1090 1095 1100
- Leu Thr Val Leu Ala Val Asp Arg Gly Ser Val Pro Leu Ser Ser Val 1105 1110 1115 1120
- Thr Glu Val Tyr Ile Glu Val Thr Asp Ala Asn Asp Asn Pro Pro Gln
  1125 1130 1135
- Met Ser Gln Ala Val Phe Tyr Pro Ser Ile Gln Glu Asp Ala Pro Val 1140 1145 1150
- Gly Thr Ser Val Leu Gln Leu Asp Ala Trp Asp Pro Asp Ser Ser Ser 1155 1160 1165
- Lys Gly Lys Leu Thr Phe Asn Ile Thr Ser Gly Asn Tyr Met Gly Phe 1170 1175 1180
- Phe Met Ile His Pro Val Thr Gly Leu Leu Ser Thr Ala Gln Gln Leu 1185 1190 1195 1200
- Asp Arg Glu Asn Lys Asp Glu His Ile Leu Glu Val Thr Val Leu Asp 1205 1210 1215
- Asn Gly Glu Pro Ser Leu Lys Ser Thr Ser Arg Val Val Val Gly Ile 1220 1225 1230
- Leu Asp Val Asn Asp Asn Pro Pro Ile Phe Ser His Lys Leu Phe Asn 1235 1240 1245
- Val Arg Leu Pro Glu Arg Leu Ser Pro Val Ser Pro Gly Pro Val Tyr 1250 1255 1260
- Arg Leu Val Ala Ser Asp Leu Asp Glu Gly Leu Asn Gly Arg Val Thr 1265 1270 1275 1280
- Tyr Ser Ile Glu Asp Ser Tyr Glu Glu Ala Phe Ser Ile Asp Leu Val 1285 1290 1295
- Thr Gly Val Val Ser Ser Asn Ser Thr Phe Thr Ala Gly Glu Tyr Asn 1300 1305 1310
- Ile Leu Thr Ile Lys Ala Thr Asp Ser Gly Gln Pro Pro Leu Ser Ala 1315 1320 1325
- Ser Val Arg Leu His Ile Glu Trp Ile Pro Trp Pro Arg Pro Ser Ser 1330 1340
- Ile Pro Leu Ala Phe Asp Glu Thr Tyr Tyr Ser Phe Thr Val Met Glu 1345 1350 1355 1360
- Thr Asp Pro Val Asn His Met Val Gly Val Ile Ser Val Glu Gly Arg 1365 1370 1375
- Pro Gly Leu Phe Trp Phe Asn Ile Ser Gly Gly Asp Lys Asp Met Asp 1380 1385 1390

- Phe Asp Ile Glu Lys Thr Thr Gly Ser Ile Val Ile Ala Arg Pro Leu 1395 1400 1405
- Asp Thr Arg Arg Ser Asn Tyr Asn Leu Thr Val Glu Val Thr Asp 1410 1415 1420
- Gly Ser Arg Thr Ile Ala Thr Gln Val His Ile Phe Met Ile Ala Asn 1425 1430 1435 1440
- Ile Asn His His Arg Pro Gln Phe Leu Glu Thr Arg Tyr Glu Val Arg 1445 1450 1455
- Val Pro Gln Asp Thr Val Pro Gly Val Glu Leu Leu Arg Val Gln Ala 1460 1465 1470
- Ile Asp Gln Asp Lys Gly Lys Ser Leu Ile Tyr Thr Ile His Gly Ser 1475 1480 1485
- Gln Asp Pro Gly Ser Ala Ser Leu Phe Gln Leu Asp Pro Ser Ser Gly 1490 1495 1500
- Val Leu Val Thr Val Gly Lys Leu Asp Leu Gly Ser Gly Pro Ser Gln 1505 1510 1515 1520
- His Thr Leu Thr Val Met Val Arg Asp Gln Glu Ile Pro Ile Lys Arg 1525 1530 1535
- Asn Phe Val Trp Val Thr Ile His Val Glu Asp Gly Asn Leu His Pro 1540 1545 1550
- Pro Arg Phe Thr Gln Leu His Tyr Glu Ala Ser Val Pro Asp Thr Ile 1555 1560 1565
- Ala Pro Gly Thr Glu Leu Leu Gln Val Arg Ala Met Asp Ala Asp Arg 1570 1580
- Gly Val Asn Ala Glu Val His Tyr Ser Leu Leu Lys Gly Asn Ser Glu 1585 1590 1595 1600
- Gly Phe Phe Asn Ile Asn Ala Leu Leu Gly Ile Ile Thr Leu Ala Gln 1605 1610 1615
- Lys Leu Asp Gln Ala Asn His Ala Pro His Thr Leu Thr Val Lys Ala 1620 1625 1630
- Glu Asp Gln Gly Ser Pro Gln Trp His Asp Leu Ala Thr Val Ile Ile 1635 1640 1645
- His Val Tyr Pro Ser Asp Arg Ser Ala Pro Ile Phe Ser Lys Ser Glu 1650 1660
- Tyr Phe Val Glu Ile Pro Glu Ser Ile Pro Val Gly Ser Pro Ile Leu 1665 1670 1680
- Leu Val Ser Ala Met Ser Pro Ser Glu Val Thr Tyr Glu Leu Arg Glu 1685 1690 1695

- Gly Asn Lys Asp Gly Val Phe Ser Met Asn Ser Tyr Ser Gly Leu Ile 1700 1705 1710
- Ser Thr Gln Lys Lys Leu Asp His Glu Lys Ile Ser Ser Tyr Gln Leu 1715 1720 1725
- Lys Ile Arg Gly Ser Asn Met Ala Gly Ala Phe Thr Asp Val Met Val 1730 1735 1740
- Val Val Asp Ile Ile Asp Glu Asn Asp Asn Ala Pro Met Phe Leu Lys 1745 1750 1760
- Ser Thr Phe Val Gly Gln Ile Ser Glu Ala Ala Pro Leu Tyr Ser Met 1765 1770 1775
- Ile Met Asp Lys Asn Asn Pro Phe Val Ile His Ala Ser Asp Ser 1780 1785 1790
- Asp Lys Glu Ala Asn Ser Leu Leu Val Tyr Lys Ile Leu Glu Pro Glu 1795 1800 1805
- Ala Leu Lys Phe Phe Lys Ile Asp Pro Ser Met Gly Thr Leu Thr Ile 1810 1815 1820
- Val Ser Glu Met Asp Tyr Glu Ser Met Pro Ser Phe Gln Phe Cys Val 1825 1830 1835 1840
- Tyr Val His Asp Gln Gly Ser Pro Val Leu Phe Ala Pro Arg Pro Ala 1845 1850 1855
- Gln Val Ile Ile His Val Arg Asp Val Asn Asp Ser Pro Pro Arg Phe 1860 1865 1870
- Ser Glu Gln Ile Tyr Glu Val Ala Ile Val Gly Pro Ile His Pro Gly 1875 1880 1885
- Met Glu Leu Leu Met Val Arg Ala Ser Asp Glu Asp Ser Glu Val Asn 1890 1895 1900
- Tyr Ser Ile Lys Thr Gly Asn Ala Asp Glu Ala Val Thr Ile His Pro 1905 1910 1915 1920
- Val Thr Gly Ser Ile Ser Val Leu Asn Pro Ala Phe Leu Gly Leu Ser 1925 1930 1935
- Arg Lys Leu Thr Ile Arg Ala Ser Asp Gly Leu Tyr Gln Asp Thr Ala 1940 1945 1950
- Leu Val Lys Ile Ser Leu Thr Gln Val Leu Asp Lys Ser Leu Gln Phe 1955 1960 1965
- Asp Gln Asp Val Tyr Trp Ala Ala Val Lys Glu Asn Leu Gln Asp Arg 1970 1975 1980
- Lys Ala Leu Val Ile Leu Gly Ala Gln Gly Asn His Leu Asn Asp Thr 1985 1990 1995 2000

- Leu Ser Tyr Phe Leu Leu Asn Gly Thr Asp Met Phe His Met Val Gln 2005 2010 2015
- Ser Ala Gly Val Leu Gln Thr Arg Gly Val Ala Phe Asp Arg Glu Gln 2020 2025 2030
- Gln Asp Thr His Glu Leu Ala Val Glu Val Arg Asp Asn Arg Thr Pro 2035 2040 2045
- Gln Arg Val Ala Gln Gly Leu Val Arg Val Ser Ile Glu Asp Val Asn 2050 2055 2060
- Asp Asn Pro Pro Lys Phe Lys His Leu Pro Tyr Tyr Thr Ile Ile Gln 2065 2070 2075 2080
- Asp Gly Thr Glu Pro Gly Asp Val Leu Phe Gln Val Ser Ala Thr Asp 2085 2090 2095
- Glu Asp Leu Gly Thr Asn Gly Ala Val Thr Tyr Glu Phe Ala Glu Asp 2100 2105 2110
- Tyr Thr Tyr Phe Arg Ile Asp Pro Tyr Leu Gly Asp Ile Ser Leu Lys 2115 2120 2125
- Lys Pro Phe Asp Tyr Gln Ala Leu Asn Lys Tyr His Leu Lys Val Ile 2130 2135 2140
- Ala Arg Asp Gly Gly Thr Pro Ser Leu Gln Ser Glu Glu Glu Val Leu 2145 2150 2155 2160
- Val Thr Val Arg Asn Lys Ser Asn Pro Leu Phe Gln Ser Pro Tyr Tyr 2165 2170 2175
- Lys Val Arg Val Pro Glu Asn Ile Thr Leu Tyr Thr Pro Ile Leu His 2180 2185 2190
- Thr Gln Ala Arg Ser Pro Glu Gly Leu Arg Leu Ile Tyr Asn Ile Val 2195 2200 2205
- Glu Glu Glu Pro Leu Met Leu Phe Thr Thr Asp Phe Lys Thr Gly Val 2210 2220
- Leu Thr Val Thr Gly Pro Leu Asp Tyr Glu Ser Lys Thr Lys His Val 2225 2230 2235 2240
- Phe Thr Val Arg Ala Thr Asp Thr Ala Leu Gly Ser Phe Ser Glu Ala 2245 2250 2255
- Thr Val Glu Val Leu Val Glu Asp Val Asn Asp Asn Pro Pro Thr Phe 2260 2265 2270
- Ser Gln Leu Val Tyr Thr Thr Ser Ile Ser Glu Gly Leu Pro Ala Gln 2275 2280 2285
- Thr Pro Val Ile Gln Leu Leu Ala Ser Asp Gln Asp Ser Gly Arg Asn 2290 2295 2300

- Arg Asp Val Ser Tyr Gln Ile Val Glu Asp Gly Ser Asp Val Ser Lys 2305 2310 2315 2320
- Phe Phe Gln Ile Asn Gly Ser Thr Gly Glu Met Ser Thr Val Gln Glu 2325 2330 2335
- Leu Asp Tyr Glu Ala Gln Gln His Phe His Val Lys Val Arg Ala Met 2340 2345 2350
- Asp Lys Gly Asp Pro Pro Leu Thr Gly Glu Thr Leu Val Val Asn 2355 2360 2365
- Val Ser Asp Ile Asn Asp Asn Pro Pro Glu Phe Arg Gln Pro Gln Tyr 2370 2375 2380
- Glu Ala Asn Val Ser Glu Leu Ala Thr Cys Gly His Leu Val Leu Lys 2385 2390 2395 2400
- Val Gln Ala Ile Asp Pro Asp Ser Arg Asp Thr Ser Arg Leu Glu Tyr 2405 2410 2415
- Leu Ile Leu Ser Gly Asn Gln Asp Arg His Phe Phe Ile Asn Ser Ser 2420 2425 2430
- Ser Gly Ile Ile Ser Met Phe Asn Leu Cys Lys Lys His Leu Asp Ser 2435 2440 2445
- Ser Tyr Asn Leu Arg Val Gly Ala Ser Asp Gly Val Phe Arg Ala Thr 2450 2455 2460
- Val Pro Val Tyr Ile Asn Thr Thr Asn Ala Asn Lys Tyr Ser Pro Glu 2465 2470 2475 2480
- Phe Gln Gln His Leu Tyr Glu Ala Glu Leu Ala Glu Asn Ala Met Val 2485 2490 2495
- Gly Thr Lys Val Ile Asp Leu Leu Ala Ile Asp Lys Asp Ser Gly Pro 2500 2505 2510
- Tyr Gly Thr Ile Asp Tyr Thr Ile Ile Asn Lys Leu Ala Ser Glu Lys 2515 2520 2525
- Phe Ser Ile Asn Pro Asn Gly Gln Ile Ala Thr Leu Gln Lys Leu Asp 2530 2540
- Arg Glu Asn Ser Thr Glu Arg Val Ile Ala Ile Lys Val Met Ala Arg 2545 2550 2560
- Asp Gly Gly Gly Arg Val Ala Phe Cys Thr Val Lys Ile Ile Leu Thr 2565 2570 2575
- Asp Glu Asn Asp Asn Pro Pro Gln Phe Lys Ala Ser Glu Tyr Thr Val 2580 2585 2590
- Ser Ile Gln Ser Asn Val Ser Lys Asp Ser Pro Val Ile Gln Val Leu 2595 2600 2605

- Ala Tyr Asp Ala Asp Glu Gly Gln Asn Ala Asp Val Thr Tyr Ser Val 2610 2615 2620
- Asn Pro Glu Asp Leu Val Lys Asp Val Ile Glu Ile Asn Pro Val Thr 2625 2630 2635 2640
- Gly Val Val Lys Val Lys Asp Ser Leu Val Gly Leu Glu Asn Gln Thr 2645 2650 2655
- Leu Asp Phe Phe Ile Lys Ala Gln Asp Gly Gly Pro Pro His Trp Asn 2660 2665 2670
- Ser Leu Val Pro Val Arg Leu Gln Val Val Pro Lys Lys Val Ser Leu 2675 2680 2685
- Pro Lys Phe Ser Glu Pro Leu Tyr Thr Phe Ser Ala Pro Glu Asp Leu 2690 2695 2700
- Pro Glu Gly Ser Glu Ile Gly Ile Val Lys Ala Val Ala Ala Gln Asp 2705 2710 2715 2720
- Pro Val Ile Tyr Ser Leu Val Arg Gly Thr Thr Pro Glu Ser Asn Lys 2725 2730 2735
- Asp Gly Val Phe Ser Leu Asp Pro Asp Thr Gly Val Ile Lys Val Arg 2740 2745 2750
- Lys Pro Met Asp His Glu Ser Thr Lys Leu Tyr Gln Ile Asp Val Met 2755 2760 2765
- Ala His Cys Leu Gln Asn Thr Asp Val Val Ser Leu Val Ser Val Asn 2770 2785
- Ile Gln Val Gly Asp Val Asn Asp Asn Arg Pro Val Phe Glu Ala Asp 2785 2790 2795 2800
- Pro Tyr Lys Ala Val Leu Thr Glu Asn Met Pro Val Gly Thr Ser Val 2805 2810 2815
- Ile Gln Val Thr Ala Ile Asp Lys Asp Thr Gly Arg Asp Gly Gln Val 2820 2825 2830
- Ser Tyr Arg Leu Ser Ala Asp Pro Gly Ser Asn Val His Glu Leu Phe 2835 2840 2845
- Ala Ile Asp Ser Glu Ser Gly Trp Ile Thr Thr Leu Gln Glu Leu Asp 2850 2855 2860
- Cys Glu Thr Cys Gln Thr Tyr His Phe His Val Val Ala Tyr Asp His 2865 2870 2875 2880
- Gly Gln Thr Ile Gln Leu Ser Ser Gln Ala Leu Val Gln Val Ser Ile 2885 2890 2895
- Thr Asp Glu Asn Asp Asn Ala Pro Arg Phe Ala Ser Glu Glu Tyr Arg 2900 2905 2910

- Gly Ser Val Val Glu Asn Ser Glu Pro Gly Glu Leu Val Ala Thr Leu 2915 2920 2925
- Lys Thr Leu Asp Ala Asp Ile Ser Glu Gln Asn Arg Gln Val Thr Cys 2930 2935 2940
- Tyr Ile Thr Glu Gly Asp Pro Leu Gly Gln Phe Gly Ile Ser Gln Val 2945 2950 2955 2960
- Gly Asp Glu Trp Arg Ile Ser Ser Arg Lys Thr Leu Asp Arg Glu His
  2965 2970 2975
- Thr Ala Lys Tyr Leu Leu Arg Val Thr Ala Ser Asp Gly Lys Phe Gln 2980 2985 2990
- Ala Ser Val Thr Val Glu Ile Phe Val Leu Asp Val Asn Asp Asn Ser 2995 3000 3005
- Pro Gln Cys Ser Gln Leu Leu Tyr Thr Gly Lys Val His Glu Asp Val 3010 3015 3020
- Phe Pro Gly His Phe Ile Leu Lys Val Ser Ala Thr Asp Leu Asp Thr 3025 3030 3035 3040
- Asp Thr Asn Ala Gln Ile Thr Tyr Ser Leu His Gly Pro Gly Ala His 3045 3050 3055
- Glu Phe Lys Leu Asp Pro His Thr Gly Glu Leu Thr Thr Leu Thr Ala 3060 3065 3070
- Leu Asp Arg Glu Arg Lys Asp Val Phe Asn Leu Val Ala Lys Ala Thr 3075 3080 3085
- Asp Gly Gly Gly Arg Ser Cys Gln Ala Asp Ile Thr Leu His Val Glu 3090 3095 3100
- Asp Val Asn Asp Asn Ala Pro Arg Phe Phe Pro Ser His Cys Ala Val 3105 3110 3120
- Ala Val Phe Asp Asn Thr Thr Val Lys Thr Pro Val Ala Val Val Phe 3125 3130 3135
- Ala Arg Asp Pro Asp Gln Gly Ala Asn Ala Gln Val Val Tyr Ser Leu 3140 3145 3150
- Pro Asp Ser Ala Glu Gly His Phe Ser Ile Asp Ala Thr Thr Gly Val 3155 3160 3165
- Ile Arg Leu Glu Lys Pro Leu Gln Val Arg Pro Gln Ala Pro Leu Glu 3170 3180
- Leu Thr Val Arg Ala Ser Asp Leu Gly Thr Pro Ile Pro Leu Ser Thr 3185 3190 3195 3200
- Leu Gly Thr Val Thr Val Ser Val Val Gly Leu Glu Asp Tyr Leu Pro
  3205 3210 3215

- Val Phe Leu Asn Thr Glu His Ser Val Gln Val Pro Glu Asp Ala Pro 3220 3225 3230
- Pro Gly Thr Glu Val Leu Gln Leu Ala Thr Leu Thr Arg Pro Gly Ala 3235 3240 3245
- Glu Lys Thr Gly Tyr Arg Val Val Ser Gly Asn Glu Gln Gly Arg Phe 3250 3255 3260
- Arg Leu Asp Ala Arg Thr Gly Ile Leu Tyr Val Asn Ala Ser Leu Asp 3265 3270 3275 3280
- Phe Glu Thr Ser Pro Lys Tyr Phe Leu Ser Ile Glu Cys Ser Arg Lys 3285 3290 3295
- Ser Ser Ser Leu Ser Asp Val Thr Thr Val Met Val Asn Ile Thr 3300 3305 3310
- Asp Val Asn Glu His Arg Pro Gln Phe Pro Gln Asp Pro Tyr Ser Thr 3315 3320 3325
- Arg Val Leu Glu Asn Ala Leu Val Gly Asp Val Ile Leu Thr Val Ser 3330 3335 3340
- Ala Thr Asp Glu Asp Gly Pro Leu Asn Ser Asp Ile Thr Tyr Ser Leu 3345 3350 3360
- Ile Gly Gly Asn Gln Leu Gly His Phe Thr Ile His Pro Lys Lys Gly 3365 3370 3375
- Glu Leu Gln Val Ala Lys Ala Leu Asp Arg Glu Gln Ala Ser Ser Tyr 3380 3385 3390
- Ser Leu Lys Leu Arg Ala Thr Asp Ser Gly Gln Pro Pro Leu His Glu 3395 3400 3405
- Asp Thr Asp Ile Ala Ile Gln Val Ala Asp Val Asn Asp Asn Pro Pro 3410 3415 3420
- Arg Phe Phe Gln Leu Asn Tyr Ser Thr Thr Val Gln Glu Asn Ser Pro 3425 3430 3435 3440
- Ile Gly Ser Lys Val Leu Gln Leu Ile Leu Ser Asp Pro Asp Ser Pro 3445 3450 3455
- Glu Asn Gly Pro Pro Tyr Ser Phe Arg Ile Thr Lys Gly Asn Asn Gly 3460 3465 3470
- Ser Ala Phe Arg Val Thr Pro Asp Gly Trp Leu Val Thr Ala Glu Gly 3475 3480 3485
- Leu Ser Arg Arg Ala Gln Glu Trp Tyr Gln Leu Gln Ile Gln Ala Ser 3490 3495 3500
- Asp Ser Gly Ile Pro Pro Leu Ser Ser Leu Thr Ser Val Arg Val His 3505 3510 3515 3520

- Val Thr Glu Gln Ser His Tyr Ala Pro Ser Ala Leu Pro Leu Glu Ile 3525 3530 3535
- Phe Ile Thr Val Gly Glu Asp Glu Phe Gln Gly Gly Met Val Gly Lys 3540 3545 3550
- Ile His Ala Thr Asp Arg Asp Pro Gln Asp Thr Leu Thr Tyr Ser Leu 3555 3560 3565
- Ala Glu Glu Glu Thr Leu Gly Arg His Phe Ser Val Gly Ala Pro Asp 3570 3575 3580
- Gly Lys Ile Ile Ala Ala Gln Gly Leu Pro Arg Gly His Tyr Ser Phe 3585 3590 3595 3600
- Asn Val Thr Val Ser Asp Gly Thr Phe Thr Thr Ala Gly Val His 3605 3610 3615
- Val Tyr Val Trp His Val Gly Gln Glu Ala Leu Gln Gln Ala Met Trp 3620 3635 3630
- Met Gly Phe Tyr Gln Leu Thr Pro Glu Glu Leu Val Ser Asp His Trp 3635 3640 3645
- Arg Asn Leu Gln Arg Phe Leu Ser His Lys Leu Asp Ile Lys Arg Ala 3650 3655 3660
- Asn Ile His Leu Ala Ser Leu Gln Pro Ala Glu Ala Val Ala Gly Val 3665 3670 3680
- Asp Val Leu Leu Val Phe Glu Gly His Ser Gly Thr Phe Tyr Glu Phe 3685 3690 3695
- Gln Glu Leu Ala Ser Ile Ile Thr His Ser Ala Lys Glu Met Glu His 3700 3705 3710
- Ser Val Gly Val Gln Met Arg Ser Ala Met Pro Met Val Pro Cys Gln 3715 3720 3725
- Gly Pro Thr Cys Gln Gly Gln Ile Cys His Asn Thr Val His Leu Asp 3730 3735 3740
- Pro Lys Val Gly Pro Thr Tyr Ser Thr Ala Arg Leu Ser Ile Leu Thr 3745 3750 3755 3760
- Pro Arg His His Leu Gln Arg Ser Cys Ser Cys Asn Gly Thr Ala Thr 3765 3770 3775
- Arg Phe Ser Gly Gln Ser Tyr Val Arg Tyr Arg Ala Pro Ala Ala Arg 3780 3785 3790
- Asn Trp His Ile His Phe Tyr Leu Lys Thr Leu Gln Pro Gln Ala Ile 3795 3800 3805
- Leu Leu Phe Thr Asn Glu Thr Ala Ser Val Ser Leu Lys Leu Ala Ser 3810 3820

- Gly Val Pro Gln Leu Glu Tyr His Cys Leu Gly Gly Phe Tyr Gly Asn 3825 3830 3835 3840
- Leu Ser Ser Gln Arg His Val Asn Asp His Glu Trp His Ser Ile Leu 3845 3850 3855
- Val Glu Glu Met Asp Ala Ser Ile Arg Leu Met Val Asp Ser Met Gly 3860 3865 3870
- Asn Thr Ser Leu Val Val Pro Glu Asn Cys Arg Gly Leu Arg Pro Glu 3875 3880 3885
- Arg His Leu Leu Gly Gly Leu Ile Leu Leu His Ser Ser Ser Asn 3890 3895 3900
- Val Ser Gln Gly Phe Glu Gly Cys Leu Asp Ala Val Val Asn Glu 3905 3910 3915 3920
- Glu Ala Leu Asp Leu Leu Ala Pro Gly Lys Thr Val Ala Gly Leu Leu 3925 3930 3935
- Glu Thr Gln Ala Leu Thr Gln Cys Cys Leu His Ser Asp Tyr Cys Ser 3940 3945 3950
- Gln Asn Thr Cys Leu Asn Gly Gly Lys Cys Ser Trp Thr His Gly Ala 3955 3960 3965
- Gly Tyr Val Cys Lys Cys Pro Pro Gln Phe Ser Gly Lys His Cys Glu 3970 3975 3980
- Gln Gly Arg Glu Asn Cys Thr Phe Ala Pro Cys Leu Glu Gly Gly Thr 3985 3990 3995 4000
- Cys Ile Leu Ser Pro Lys Gly Ala Ser Cys Asn Cys Pro His Pro Tyr 4005 4010 4015
- Thr Gly Asp Arg Cys Glu Met Glu Ala Arg Gly Cys Ser Glu Gly His
  4020 4025 4030
- Cys Leu Val Thr Pro Glu Ile Gln Arg Gly Asp Trp Gly Gln Gln Glu 4035 4040 4045
- Leu Leu Ile Ile Thr Val Ala Val Ala Phe Ile Ile Ile Ser Thr Val 4050 4055 4060
- Gly Leu Leu Phe Tyr Cys Arg Arg Cys Lys Ser His Lys Pro Val Ala 4065 4070 4075 4080
- Met Glu Asp Pro Asp Leu Leu Ala Arg Ser Val Gly Val Asp Thr Gln
  4085 4090 4095
- Ala Met Pro Ala Ile Glu Leu Asn Pro Leu Ser Ala Ser Ser Cys Asn 4100 4105 4110
- Asn Leu Asn Gln Pro Glu Pro Ser Lys Ala Ser Val Pro Asn Glu Leu
  4115 4120 4125

- Val Thr Phe Gly Pro Asn Ser Lys Gln Arg Pro Val Val Cys Ser Val 4130 4135 4140
- Pro Pro Arg Leu Pro Pro Ala Ala Val Pro Ser His Ser Asp Asn Glu 4145 4150 4155 4160
- Pro Val Ile Lys Arg Thr Trp Ser Ser Glu Glu Met Val Tyr Pro Gly
  4165 4170 4175
- Gly Ala Met Val Trp Pro Pro Thr Tyr Ser Arg Asn Glu Arg Trp Glu
  4180 4185 4190
- Tyr Pro His Ser Glu Val Thr Gln Gly Pro Leu Pro Pro Ser Ala His 4195 4200 4205
- Arg His Ser Thr Pro Val Val Met Pro Glu Pro Asn Gly Leu Tyr Gly 4210 4215 4220
- Gly Phe Pro Phe Pro Leu Glu Met Glu Asn Lys Arg Ala Pro Leu Pro 4225 4230 4235 4240
- Pro Arg Tyr Ser Asn Gln Asn Leu Glu Asp Leu Met Pro Ser Arg Pro 4245 4250 4255
- Pro Ser Pro Arg Glu Arg Leu Val Ala Pro Cys Leu Asn Glu Tyr Thr 4260 4265 4270
- Ala Ile Ser Tyr Tyr His Ser Gln Phe Arg Gln Gly Gly Gly Pro
  4275 4280 4285
- Cys Leu Ala Asp Gly Gly Tyr Lys Gly Val Gly Met Arg Leu Ser Arg 4290 4295 4300
- Ala Gly Pro Ser Tyr Ala Val Cys Glu Val Glu Gly Ala Pro Leu Ala 4305 4310 4315 4320
- Gly Gln Gly Gln Pro Arg Val Pro Pro Asn Tyr Glu Gly Ser Asp Met 4325 4330 4335
- Val Glu Ser Asp Tyr Gly Ser Cys Glu Glu Val Met Phe
  4340
  4345

<210> 20

<211> 4351

<212> PRT

<213> Rattus norvegicus

<400> 20

- Met Thr Leu Val Leu Leu Gly Leu Ala Ile Leu Leu Leu His Arg Ala 1 5 10 15
- Ala Cys Glu Lys Ser Leu Glu Glu Thr Ile Pro Pro Leu Ser Trp Arg
  20 25 30
- Phe Thr His Ser Leu Tyr Asn Ala Thr Ile Tyr Glu Asn Ser Ala Pro 35 40 45

Lys Thr Tyr Val Glu Ser Pro Val Lys Met Gly Met Tyr Leu Ala Glu 55 Pro His Trp Val Val Lys Tyr Arg Ile Ile Ser Gly Asp Ala Ala Gly Val Phe Lys Thr Glu Glu His Val Val Gly Asn Phe Cys Phe Leu Arg Ile Arg Thr Lys Ser Ser Asn Thr Ala Leu Leu Asn Arg Glu Val Arg 105 Asp Ser Tyr Thr Leu Ile Val Gln Ala Ser Asp Lys Ser Leu Glu Phe 120 Glu Ala Leu Thr Gln Val Val Val His Ile Leu Asp Gln Asn Asp Leu 135 Lys Pro Leu Phe Ser Pro Pro Ser Tyr Arg Val Thr Ile Ser Glu Asp 150 155 Arg Pro Leu Lys Ser Pro Ile Cys Lys Val Thr Ala Thr Asp Ala Asp Leu Gly Gln Asn Ala Glu Phe Tyr Tyr Ala Phe Asn Ala Arg Ser Glu 185 Val Phe Ala Ile His Pro Thr Ser Gly Val Val Thr Val Ala Gly Lys 200 Leu Asn Val Thr Arg Arg Gly Lys Tyr Glu Leu Gln Val Leu Ala Val Asp Arg Met Arg Lys Ile Ser Glu Gly Asn Gly Phe Gly Asn Leu Ala 230 235 Ser Leu Val Ile Arg Val Glu Pro Val His Arg Lys Pro Pro Ala Ile 245 Asn Leu Val Val Leu Asn Pro Pro Glu Gly Asp Glu Gly Asp Ile Tyr Ala Ile Val Thr Val Asp Thr Asn Gly Ser Gly Ala Glu Val Asp Ser 275 Leu Glu Val Val Gly Gly Asp Pro Gly Lys Tyr Phe Lys Val Leu Arg Ser Tyr Ala Gln Gly Asn Glu Phe Asn Leu Val Ala Val Arg Asp Ile 310 315 Asn Trp Ala Glu His Pro His Gly Phe Asn Ile Ser Leu Gln Thr His 325 Ser Trp Ser Arg Phe Pro Pro His Ser Ile Ile Arg Ala Phe His Leu 340 345 350

Pro	Ser	Trp 355	Lys	Leu	Ala	Asn	Leu 360	Arg	Phe	Glu	Lys	Ala 365	Val	Tyr	Arg
Val	Lys 370	Leu	Ser	Glu	Phe	Ser 375	Pro	Pro	Gly	Ser	Arg 380	Val	Ala	Leu	Val
Lys 385	Val	Thr	Thr	Ala	Leu 390	Pro	Asn	Leu	Arg	Tyr 395	Ser	Leu	Lys	Pro	Ser 400
Ser	Arg	Asn	Thr	Ala 405	Phe	Lys	Leu	Asn	Ala 410	Arg	Thr	Gly	Leu	Ile 415	Thr
Thr	Thr	Lys	Leu 420	Val	Asp	Phe	His	Glu 425	Gln	Asn	Gln	Tyr	Gln 430	Leu	His
Val	Lys	Thr 435	Ser	Leu	Gly	Gln	Ala 440	Thr	Thr	Thr	Val	Ile 445	Ile	Asp	Ile
Val	Asp 450	Cys	Asn	Asn	His	Ala 455	Pro	Val	Phe	Asn	Arg 460	Ser	Ser	Tyr	Glu
Gly 465	Thr	Leu	Asp	Glu	Asn 470	Ile	Pro	Pro	Gly	Thr 475	Ser	Val	Leu	Thr	Val 480
Thr	Ala	Thr	Asp	Gln 485	Asp	His	Gly	Asp	Asn 490	Gly	His	Ile	Thr	Tyr 495	Ser
Ile	Ala	Gly	Pro 500	Lys	Ala	Val	Pro	Phe 505	Ser	Ile	Asp	Pro	Leu 510	Leu	Gly
Val	Ile	Ser 515	Thr	Thr	Lys	Pro	Met 520	Asp	Tyr	Glu	Leu	Met 525	Lys	Arg	Ile
Tyr	Thr 530	Phe	Arg	Val	Arg	Ala 535	Ser	Asp	Trp	Gly	Ser 540	Pro	Phe	Arg	Gln
Glu 545	Lys	Glu	Val	Ser	Val 550	Ser	Leu	Arg	Leu	Lys 555	Asn	Leu	Asn	Asp	Asn 560
Gln	Pro	Met	Phe	Glu 565	Glu	Val	Asn	Cys	Thr 570	Val	Ser	Leu	Arg	Gln 575	Asp
Val	Pro	Val	Gly 580	Lys	Ser	Ile	Met	Ala 585	Val	Ser	Ala	Ile	Asp 590	Met	Asp
Glu	Leu	Gln 595	Asn	Leu	Lys	Tyr	Glu 600	Ile	Val	Ser	Gly	Asn 605	Glu	Gln	Asp
Tyr	Phe 610	His	Leu	Asn	His	Phe 615	Ser	Gly	Val	Ile	Ser 620	Leu	Lys	Arg	Ser
Phe 625	Met	Asn	Leu	Thr	Ala 630	Val	Arg	Pro	Thr	Ile 635	Tyr	Ser	Leu	Lys	Ile 640
Thr	Ala	Ser	Asp	Gly 645	Lys	Asn	Tyr	Ala	Ser 650	Pro	Thr	Thr	Leu	Lys 655	Val

Thr Val Val Lys Asp Pro His Ser Glu Val Pro Val Gln Cys Asp Lys 665 Thr Gly Val Leu Thr His Ile Thr Lys Thr Ile Leu Gln Ser Ala Gly 680 Leu Gln Ser Gln Glu Leu Gly Glu Glu Glu Phe Thr Ser Leu Ser Asn 695 Tyr Gln Ile Asn His His Ser Pro Gln Phe Glu Asp His Phe Pro Gln 710 715 Ser Ile Asp Ile Leu Glu Gln Val Pro Ile Asn Thr Pro Leu Ala Arg 725 730 Leu Ala Ala Thr Asp Pro Asp Thr Gly Phe His Gly Lys Leu Val Tyr 740 745 Val Ile Ser Asp Gly Asn Glu Glu Gly Cys Phe Asp Ile Glu Leu Glu Thr Gly Leu Leu Met Val Ala Ala Leu Asp Tyr Glu Thr Thr Ser 770 775 Phe Tyr Val Leu Asn Val Thr Val Tyr Asp Leu Gly Thr Pro Pro Lys 790 Ser Ser Trp Lys Leu Leu Thr Val Thr Val Lys Asp Trp Asn Asp Asn 810 Pro Pro Arg Phe Pro Pro Gly Gly Tyr Gln Leu Thr Ile Ser Glu Asp 820 Thr Glu Val Gly Thr Thr Ile Ala Glu Leu Lys Thr Glu Asp Ala Asp 840 Ser Glu Asp Asn Arg Arg Val Arg Tyr Thr Leu Leu Thr Pro Thr Glu 855 850 Lys Phe Ser Leu His Pro Phe Thr Gly Glu Leu Val Val Thr Gly His Leu Asp Arg Glu Ser Glu Ser Gln Tyr Ile Leu Lys Ala Glu Ala Arg 890 Asp Gln Pro Thr Lys Gly His Gln Leu Phe Ser Val Thr Asp Leu Ile 900 905 Val Thr Leu Glu Asp Ile Asn Asp Asn Pro Pro Gln Cys Ile Thr Glu 920 His Arg Arg Leu Lys Val Pro Glu Asp Met Pro Leu Gly Thr Val Leu 930 935 Thr Phe Leu Asp Ala Ser Asp Pro Asp Leu Gly Pro Ala Gly Glu Val 950 955 960 945

- Lys Tyr Ile Leu Val Glu Asp Ala His Gly Thr Phe Gln Val His Pro 965 970 975
- Met Thr Gly Ala Leu Ser Leu Glu Lys Glu Leu Asp Phe Glu Arg Arg 980 985 990
- Ala Gly Tyr Asn Leu Ser Phe Trp Ala Ser Asp Ser Gly Lys Pro Leu
  995 1000 1005
- Ser Arg Arg Thr Leu Cys His Val Glu Val Leu Val Met Asp Val Asn 1010 1025
- Glu Asn Leu His Ser Pro His Phe Ser Ser Phe Val Tyr Gln Gly Gln 1025 1030 1035 1040
- Val Gln Glu Asn Ser Pro Ala Gly Thr Pro Val Met Val Val Thr Ala 1045 1050 1055
- Gln Asp Asp Ser Gly Leu Asp Gly Glu Leu Gln Tyr Phe Leu Arg 1060 1065 1070
- Ala Gly Thr Gly Leu Glu Thr Phe Ser Ile Asn Gln Asp Thr Gly Met 1075 1080 1085
- Leu Glu Thr Leu Ala Pro Leu Asp Arg Glu Phe Thr Pro Tyr Tyr Trp 1090 1095 1100
- Leu Thr Val Leu Ala Val Asp Arg Gly Ser Val Pro Leu Ser Ala Val 1105 1110 1115 1120
- Thr Glu Val Tyr Ile Glu Val Thr Asp Ile Asn Asp Asn Ile Pro Ser 1125 1130 1135
- Met Ser Arg Pro Val Phe Tyr Pro Ser Val Leu Glu Asp Ala Pro Leu 1140 1145 1150
- Gly Thr Ser Val Leu Gln Leu Glu Ala Trp Asp Pro Asp Ser Ser Ser 1155 1160 1165
- Gln Gly Lys Leu Thr Phe Asn Leu Thr Ser Gly Asn His Leu Gly His 1170 1175 1180
- Phe Ile Val His Pro Phe Thr Gly Leu Leu Thr Thr Ala Lys Gln Leu 1185 1190 1195 1200
- Asp Arg Glu Asn Lys Asp Glu Tyr Val Leu Glu Val Thr Val Gln Asp 1205 1210 1215
- Asn Gly Asp Pro Ser Leu Arg Ser Thr Ser Arg Val Val Cys Ile 1220 1225 1230
- Leu Asp Val Asn Asp Asn Pro Pro Met Phe Ser His Lys Leu Phe Asn 1235 1240 1245
- Val Arg Leu Ser Glu Arg Leu Ser Pro Leu Ser Pro Glu Pro Val Tyr 1250 1255 1260

- Arg Leu Val Ala Ser Asp Pro Asp Glu Gly Leu Asn Gly Ser Val Thr 1265 1270 1275 1280
- Tyr Ser Ile Glu Glu Ser Asp Glu Glu Ser Phe Arg Ile Asp Pro Val 1285 1290 1295
- Thr Gly Val Val Ser Ser Ser Ser Thr Phe Ala Ala Gly Glu Tyr Asn 1300 1305 1310
- Ile Leu Thr Ile Lys Ala Thr Asp Ser Gly Gln Pro Ala Leu Ser Thr 1315 1320 1325
- Ser Val Arg Leu His Ile Glu Trp Ile Pro Gln Pro Arg Pro Ser Ser 1330 1340
- Ile Pro Leu Ser Phe Asp Glu Ser Tyr Tyr Ser Phe Thr Val Met Glu 1345 1350 1355 1360
- Thr Asp Pro Val Asn His Met Val Gly Val Ile Ser Val Glu Gly Arg
  1365 1370 1375
- Pro Gly Leu Phe Trp Phe His Ile Ser Asp Gly Asp Lys Asp Met Asp 1380 1385 1390
- Phe Asp Ile Glu Lys Thr Thr Gly Ser Ile Val Ile Ala Arg Pro Leu 1395 1400 1405
- Asp Thr Arg Arg Lys Ser Ser Tyr Asn Leu Thr Val Glu Val Thr Asp 1410 1415 1420
- Gly Phe His Thr Ile Ala Thr Gln Val His Ile Phe Met Ile Ala Asn 1425 1430 1435 1440
- Ile Asn His His Arg Pro Gln Phe Leu Gln Asp His Tyr Glu Ile Arg 1445 1450 1455
- Val Pro Gln Asp Thr Leu Pro Gly Val Glu Leu Leu Arg Val Gln Ala 1460 1465 1470
- Thr Asp Gln Asp His Gly Lys Gly Leu Ile Tyr Thr Ile Leu Ser Ser 1475 1480 1485
- Gln Asp Pro Gly Ser Ala Asn Leu Phe Gln Leu Asp Pro Ser Ser Gly 1490 1495 1500
- Val Leu Val Thr Val Gly Thr Leu Glu Leu His Ser Gly Pro Ser Gln 1505 1510 1515 1520
- His Ile Leu Thr Val Met Val Arg Asp Gln Glu Met Pro Ile Lys Arg 1525 1530 1535
- Asn Phe Val Trp Val Thr Ile His Val Glu Asp Gly Asn Leu His Ser 1540 1545 1550
- Pro His Phe Thr Gln Leu Arg Tyr Glu Ala Asn Val Pro Asp Thr Thr 1555 1560 1565

- Ala Pro Gly Thr Glu Leu Leu Gln Val Arg Ala Val Asp Ala Asp Arg 1570 1575 1580
- Gly Ala Asn Ala Glu Ile His Tyr Ser Phe Leu Lys Gly Asn Ser Asp 1585 1590 1595 1600
- Gly Phe Phe Asn Ile Asp Ser Leu Leu Gly Ile Ile Thr Val Ala Gln
  1605 1610 1615
- Arg Leu Tyr His Val His Leu Thr Arg His Ala Leu Thr Val Lys Ala 1620 1625 1630
- Glu Asp Gln Gly Ser Pro Arg Arg His Asp Leu Ala Leu Val Val Ile 1635 1640 1645
- His Val His Pro Ser Asp Ser Ser Ala Pro Val Phe Ser Lys Asp Glu 1650 1660
- Tyr Phe Ile Glu Ile Pro Glu Ser Val Pro Ile Gly Ser Pro Ile Leu 1665 1670 1680
- Leu Leu Ser Ala Gly Ser Ser Ser Glu Val Thr Tyr Glu Leu Arg Glu 1685 1690 1695
- Gly Asn Lys Asp Ser Val Phe Ser Met Asn Ser Tyr Ser Gly Leu Ile 1700 1705 1710
- Ser Thr Gln Lys Arg Leu Asp His Glu Lys Val Pro Ser Tyr Arg Leu 1715 1720 1725
- Arg Ile Arg Gly Ser Asn Met Ala Gly Val Phe Thr Glu Val Val Ala 1730 1735 1740
- Leu Val Tyr Ile Ile Asp Glu Asn Asp Asn Pro Pro Ala Phe Gly Lys 1745 1750 1755 1760
- Pro Thr Phe Leu Gly His Ile Ser Glu Ala Ala Pro Leu His Ser Leu 1765 1770 1775
- Ile Leu Gly Glu Asp Asn Ser Pro Leu Val Val Arg Ala Ser Asp Ser 1780 1785 1790
- Asp Arg Glu Ala Asn Ser Leu Leu Val Tyr Lys Ile Leu Glu Pro Glu 1795 1800 1805
- Ala Leu Lys Phe Phe Lys Ile Asp Pro Ser Met Gly Thr Leu Thr Thr 1810 1815 1820
- Thr Ser Glu Leu Asp Phe Glu Asp Thr Pro Leu Phe Gln Phe Asn Ile 1825 1830 1835 1840
- Tyr Val His Asp Gln Gly Thr Pro Ile Leu Phe Ala Pro Arg Ser Ala 1845 1850 1855
- Lys Val Ile Ile His Val Arg Asp Val Asn Asp Ser Pro Pro Arg Phe
  1860 1865 1870

- Ser Glu Gln Ile Tyr Glu Val Ala Val Val Glu Pro Ile His Pro Gly 1875 1880 1885
- Met Gly Leu Leu Thr Val Gln Ala Glu Asp Asn Asp Ser Arg Val Thr 1890 1895 1900
- Tyr Ser Ile Lys Thr Ser Asn Ala Asp Glu Ala Val Thr Ile His Pro 1905 1910 1915 1920
- Thr Thr Gly Gln Ile Ser Val Val Asn Pro Ala Thr Leu Arg Leu Phe
  1925 1930 1935
- Gln Lys Phe Ser Ile Arg Ala Ser Asp Gly Leu Tyr His Asp Thr Ala 1940 1945 1950
- Val Val Lys Ile Ser Leu Thr Gln Val Leu Asp Lys Ser Leu Gln Phe 1955 1960 1965
- Asp Gln Asp Val Tyr Arg Ala Arg Val Thr Glu Asn Thr Pro His Arg 1970 1975 1980
- Lys Ala Leu Val Ile Leu Gly Val His Gly Asn His Leu Asn Asp Thr 1985 1990 1995 2000
- Leu Ser Tyr Phe Leu Leu Asn Gly Thr Asp Leu Phe His Met Ile Glu 2005 2010 2015
- Ser Ala Gly Val Leu Gln Thr Arg Gly Gly Thr Phe Asp Arg Glu Gln 2020 2025 2030
- Gln Asp Thr His Glu Val Ala Val Glu Val Arg Asp Asn Arg Val Pro 2035 2040 2045
- Gln Arg Val Ala Gln Ala Leu Val Arg Val Ser Val Glu Asp Val Asn 2050 2055 2060
- Asp Asn Ile Pro Glu Phe Gln His Leu Pro Tyr Tyr Thr Val Ile Gln 2065 2070 2075 2080
- Asp Gly Thr Glu Pro Gly Asp Val Leu Phe Gln Val Ser Ala Thr Asp 2085 2090 2095
- Lys Asp Leu Gly Ala Asn Gly Ser Val Thr Tyr Gly Phe Ala Glu Asp 2100 2105 2110
- Tyr Ala Tyr Phe Arg Ile Asp Pro Tyr Val Gly Asp Ile Ser Leu Lys 2115 2120 2125
- Lys Pro Phe Asp Tyr Gln Ala Leu Asn Lys Tyr His Leu Arg Val Ile 2130 2135 2140
- Ala Arg Asp Ser Gly Ile Pro Pro Leu Gln Thr Glu Val Glu Val His 2145 2150 2155 2160
- Val Thr Val Arg Asn Lys Ser Asn Pro Leu Phe Gln Ser Pro Tyr Tyr 2165 2170 2175

- Lys Val Lys Val Pro Glu Asn Ile Thr Leu Tyr Thr Pro Ile Leu His 2180 2185 2190
- Thr Gln Ala Arg Ser Pro Glu Gly Leu Arg Leu Ile Tyr Asn Ile Val 2195 2200 2205
- Glu Glu Glu Pro Leu Met Leu Phe Thr Thr Asp Phe Lys Thr Gly Val 2210 2215 2220
- Leu Thr Val Thr Gly Pro Leu Asp Tyr Glu Ser Lys Asn Lys His Val 2225 2230 2235 2240
- Phe Thr Val Arg Ala Thr Asp Thr Ala Leu Gly Ser Phe Ser Glu Ala 2245 2250 2255
- Thr Val Glu Val Leu Val Glu Asp Ile Asn Asp Asn Pro Pro Thr Phe 2260 2265 2270
- Ser Gln Leu Val Tyr Thr Thr Ser Val Ser Glu Gly Ser Pro Ala Gln 2275 2280 2285
- Thr Pro Val Ile Gln Leu Leu Ala Ser Asp Gln Asp Ser Gly Gln Asn 2290 2295 2300
- Gln Asp Val Ser Tyr Gln Ile Val Glu Asp Gly Ser Asp Val Ser Lys 2305 2310 2315 2320
- Phe Phe Arg Ile Asn Gly Ser Thr Gly Glu Ile Phe Thr Ile Gln Glu 2325 2330 2335
- Leu Asp Tyr Glu Thr His Gln His Phe Arg Val Lys Val Arg Ala Met 2340 2345 2350
- Asp Lys Gly Asp Pro Pro Leu Thr Gly Glu Thr Leu Val Val Asn 2355 2360 2365
- Val Ser Asp Ile Asn Asp Asn Pro Pro Lys Phe Arg Glu Pro Gln Tyr 2370 2375 2380
- Glu Ala Asn Val Ser Glu Leu Ala Thr Cys Gly His Leu Val Leu Lys 2385 2390 2395 2400
- Val Gln Ala Leu Asp Pro Asp Ile Gly Asp Thr Ser Arg Leu Glu Tyr 2405 2410 2415
- Leu Ile Leu Ser Gly Asn Gln Asp Arg His Phe Ser Ile Asn Ser Thr
  2420 2425 2430
- Ser Gly Ile Ile Ser Met Phe Asn Leu Cys Lys Lys Gln Leu Asp Ser 2435 2440 2445
- Ser Tyr Asn Leu Arg Val Gly Ala Ser Asp Gly Val Phe Arg Ala Thr 2450 2455 2460
- Val Pro Val Tyr Ile Asn Thr Thr Asn Ala Asn Lys Tyr Ser Pro Glu 2465 2470 2475 2480

- Phe Gln Gln Asn Val Tyr Glu Ala Glu Leu Ala Glu Asn Ala Lys Val 2485 2490 2495
- Gly Thr Lys Val Ile Glu Leu Leu Ala Ile Asp Lys Asp Ser Gly Pro 2500 2505 2510
- Tyr Gly Thr Val Asp Tyr Thr Ile Ile Asn Lys Leu Ala Gly Glu Arg 2515 2520 2525
- Phe Phe Ile Asn Pro Arg Gly Gln Ile Thr Thr Leu Gln Lys Leu Asp 2530 2540
- Arg Glu Asn Ser Thr Glu Arg Val Ile Ala Ile Lys Val Met Ala Arg 2545 2550 2555 2560
- Asp Gly Gly Lys Val Ala Phe Cys Thr Val Lys Ile Ile Leu Thr 2565 2570 2575
- Asp Glu Asn Asp Asn Ala Pro Gln Phe Lys Ala Ser Gly Tyr Thr Val 2580 2585 2590
- Ser Ile Pro Ser Asn Val Ser Arg Asp Ser Pro Ile Ile Gln Val Leu 2595 2600 2605
- Ala Tyr Asp Ala Asp Glu Gly Arg Asn Ala Asp Val Thr Tyr Ser Val 2610 2620
- Asp Ser Thr Glu Asp Leu Ala Glu Glu Ile Ile Glu Val Asn Pro Thr 2625 2630 2635 2640
- Thr Gly Val Val Lys Val Lys Glu Ser Leu Val Gly Leu Glu Asn Arg 2645 2650 2655
- Ala Val Asp Phe Asn Ile Lys Ala Gln Asp Gly Gly Pro Pro His Trp 2660 2665 2670
- Asp Ser Leu Val Pro Val Arg Leu Gln Val Val Pro Asn Glu Ile Pro 2675 2680 2685
- Leu Pro Lys Phe Ser Glu Pro Leu Tyr Thr Phe Ser Ala Pro Glu Asp 2690 2695 2700
- Leu Pro Glu Gly Ser Glu Ile Gly Ser Val Lys Ala Val Ala Ala Gln 2705 2710 2715 2720
- Asp Pro Ile Ile Tyr Ser Leu Val Gln Gly Thr Thr Pro Glu Ser Asn 2725 2730 2735
- Ser Asp Asp Val Phe Ser Leu Asp Gln Asp Thr Gly Val Leu Lys Val 2740 2745 2750
- Arg Lys Ala Met Asp His Glu Ser Thr Lys Trp Tyr Gln Ile Asp Leu 2755 2760 2765
- Met Ala His Cys Pro His Glu Asp Thr Asp Leu Val Ser Leu Val Ser 2770 2775 2780

- Val Ser Ile Gln Val Glu Asp Val Asn Asp Asn Arg Pro Val Phe Glu 2785 2790 2795 2800
- Ala Asp Pro Tyr Lys Ala Phe Leu Thr Glu Asn Met Pro Gly Gly Thr 2805 2810 2815
- Thr Val Ile Gln Val Thr Ala Asn Asp Gln Asp Thr Gly Ser Asp Gly 2820 2825 2830
- Gln Val Ser Tyr Arg Leu Ser Val Glu Pro Gly Ser Asn Ile His Glu 2835 2840 2845
- Leu Phe Ala Val Asp Ser Glu Ser Gly Trp Ile Thr Thr Leu Gln Glu 2850 2855 2860
- Leu Asp Cys Glu Thr Gln Gln Thr Tyr Arg Phe Tyr Val Val Ala Phe 2865 2870 2875 2880
- Asp His Gly Gln Thr Ile Gln Leu Ser Ser Gln Ala Leu Val Glu Val 2885 2890 2895
- Ser Ile Thr Asp Glu Asn Asp Asn Pro Pro Arg Phe Ala Ser Glu Asp 2900 2905 2910
- Tyr Arg Gly Ser Val Val Glu Asn Asn Glu Pro Gly Glu Leu Val Ala 2915 2920 2925
- Thr Leu Lys Thr Leu Asp Ala Asp Val Ser Asp Gln Asn Arg Gln Val 2930 2935 2940
- Thr Cys Tyr Ile Thr Glu Gly Asp Pro Leu Gly Gln Phe Ser Ile Ser 2945 2950 2955 2960
- Gln Val Gly Asp Glu Trp Arg Ile Ser Ser Arg Lys Thr Leu Asp Arg 2965 2970 2975
- Glu His Ile Ala Lys Tyr Leu Leu Arg Val Thr Ala Ser Asp Gly Lys 2980 2985 2990
- Phe Gln Ala Ser Val Pro Val Glu Val Phe Val Val Asp Ile Asn Asp 2995 3000 3005
- Asn Ser Pro Gln Cys Ser Gln Leu Leu Tyr Thr Gly Lys Val Arg Glu 3010 3015 3020
- Asp Val Thr Pro Gly His Phe Ile Leu Lys Val Ser Ala Ile Asp Val 3025 3030 3035 3040
- Asp Met Asp Thr Asn Ala Gln Ile Thr Tyr Ser Leu His Gly Pro Gly 3045 3050 3055
- Ala Gln Glu Phe Lys Leu Asp Pro His Thr Gly Glu Leu Thr Thr Leu 3060 3065 3070
- Thr Val Leu Asp Arg Glu Arg Lys Asp Val Tyr Asn Leu Val Ala Lys 3075 3080 3085

- Ala Thr Asp Gly Gly Gln Ser Cys Gln Ala Glu Val Thr Leu His 3090 3095 3100
- Ile Glu Asp Val Asn Asp Asn Ala Pro Arg Phe Phe Pro Ser His Cys 3105 3110 3115 3120
- Asp Val Ala Val Phe Asp Asn Thr Thr Val Lys Thr Pro Val Ala Val
  3125 3130 3135
- Val Phe Ala Arg Asp Pro Asp Gln Gly Ala Asn Ala Gln Val Tyr 3140 3145 3150
- Ser Leu Thr Asp Ser Ala Asp Gly Gln Phe Ser Ile Asp Ala Thr Ser 3155 3160 3165
- Gly Val Ile Arg Leu Glu Lys Pro Leu Gln Val Arg Ala Ser Ser Ala 3170 3180
- Val Glu Leu Thr Val Arg Ala Ser Asp Leu Gly Thr Pro Ile Pro Leu 3185 3190 3195 3200
- Ser Thr Leu Gly Thr Val Thr Val Ser Val Val Gly Leu Glu Asp Tyr 3205 3210 3215
- Leu Pro Ile Phe Leu Asn Ala Glu His Ser Thr Gln Val Pro Glu Asp 3220 3225 3230
- Ala Pro Ile Asp Met Glu Val Leu His Leu Ala Thr Leu Thr Arg Pro 3235 3240 3245
- Gly Ser Glu Lys Thr Gly Tyr His Ile Thr Gly Gly Asn Glu Gln Gly 3250 3255 3260
- Lys Phe Arg Leu Asp Ala His Thr Gly Ile Leu Tyr Val Asn Gly Ser 3265 3270 3275 3280
- Leu Asp Phe Glu Thr Asn Pro Lys Tyr Phe Leu Ser Ile Glu Cys Ser 3295
- Arg Lys Ser Ser Ser Ser Leu Ser Asp Val Thr Thr Ile Val Ile Asn 3300 3305 3310
- Val Thr Asp Val Asn Glu His His Pro Arg Phe Thr His Asp Leu Tyr 3315 3320 3325
- Thr Val Arg Val Leu Glu Asn Ala Val Val Gly Asp Val Ile Leu Thr 3330 3340
- Val Ser Ala Ser Asp Asp Gly Pro Val Asn Ser Ala Ile Thr Tyr 3345 3350 3355 3360
- Ser Leu Val Gly Gly Asn Gln Leu Gly His Phe Thr Ile Asn Pro Lys 3365 3370 3375
- Lys Gly Lys Leu Gln Val Ala Lys Ala Leu Asp Trp Glu Gln Thr Pro 3380 3385 3390

- Ser Tyr Ser Leu Arg Leu Arg Ala Thr Asp Ser Gly Gln Pro Pro Leu 3395 3400 3405
- His Glu Asp Thr Glu Val Ala Val Glu Val Val Asp Val Asn Asp Asn 3410 3415 3420
- Pro Pro Arg Phe Phe Gln Leu Asn Tyr Ser Thr Ser Val Gln Glu Asn 3425 3430 3435 3440
- Ser Pro Ile Gly Ile Lys Val Leu Gln Leu Ile Leu Asp Asp Pro Asp 3445 3450 3455
- Ser Pro Gln Asn Gly Pro Pro Tyr Phe Phe Arg Ile Thr Glu Gly Asn 3460 3465 3470
- Thr Gly Ser Val Phe Arg Val Thr Pro Asp Gly Trp Leu Val Thr Ala 3475 3480 3485
- Ala Ser Leu Ser Lys Lys Ala Arg Glu Trp Tyr Gln Leu His Ile Glu 3490 3495 3500
- Val Ser Asp Ser Gly Leu Pro Pro Leu Ser Ser Ser Thr Leu Val Arg 3505 3510 3515 3520
- Val Gln Val Thr Glu Gln Ser Arg Tyr Pro Pro Ser Thr Leu Pro Leu 3525 3530 3535
- Glu Ile Ser Ile Thr Lys Gly Glu Glu Glu Phe Gln Gly Gly Met Ile 3540 3545 3550
- Gly Lys Ile His Ala Thr Asp Arg Asp Pro Gln Asp Thr Leu Thr Tyr 3555 3560 3565
- Ser Leu Glu Gln Glu Gly Gly Leu Asp Arg Tyr Phe Thr Val Gly Ala 3570 3580
- Ser Asp Gly Lys Ile Ile Ala Ser Gln Gly Leu Pro His Gly Arg Tyr 3585 3590 3595 3600
- Ser Phe Asn Val Thr Val Ser Asp Gly Thr Phe Thr Thr Thr Gly 3605 3610 3615
- Val His Val His Val Trp His Met Glu Pro Glu Val Pro Gln Gln Ala 3620 3625 3630
- Val Trp Leu Gly Phe His Gln Leu Thr Pro Glu Glu Leu Val Ser Asp 3635 3640 3645
- His Trp Arg Asn Leu Gln Arg Phe Leu Ser Asn Leu Leu Asp Val Lys 3650 3650
- Arg Ala Asn Ile His Leu Ala Ser Leu Gln Pro Ala Glu Val Thr Ala 3665 3670 3680
- Gly Val Asp Val Leu Leu Val Phe Glu Arg His Ser Gly Thr Ser Tyr 3685 3690 3695

- Asp Leu Gln Glu Leu Ala Ser Ala Ile Ala His Ser Val Arg Glu Ile 3700 3705 3710
- Glu His Ser Val Gly Ile Arg Met Arg Ser Ala Leu Pro Val Val Pro 3715 3720 3725
- Cys Gln Gly Gln Ser Cys Gln Asp Gln Thr Cys Gln Glu Thr Val Ser 3730 3740
- Leu Glu Pro Arg Val Gly Pro Ser Tyr Ser Thr Ala Arg Leu Ser Ile 3745 3750 3755 3760
- Leu Thr Pro Arg His His Leu Gly Arg Asn Cys Ser Cys Asn Gly Thr 3765 3770 3775
- Thr Leu Arg Phe Ser Gly Gln Ser Tyr Val Gln Tyr Arg Pro Leu Glu 3780 3785 3790
- Ala Gln Asn Trp Gln Ile His Phe Tyr Leu Lys Thr Leu Gln Pro Trp 3795 3800 3805
- Ala Leu Leu Met Phe Thr Asn Glu Thr Ala Ser Ile Ser Leu Lys Leu 3810 3815 3820
- Ala Asn Gly Phe Ser His Leu Glu Tyr His Cys Pro Gly Gly Phe Tyr 3825 3830 3835 3840
- Gly Asn Leu Ser Ser Arg Tyr Pro Val Asn Asp Gly Gln Trp His Ser 3845 3850 3855
- Met Leu Leu Glu Glu Arg Asp Thr Ser Val His Leu Leu Val Asp Ile 3860 3865 3870
- Thr Asp Asn Ala Ser Leu Val Ile Pro Glu Glu Cys Gln Gly Leu Arg 3875 3880 3885
- Thr Glu Arg Gln Leu Leu Gly Gly Leu Val Pro Ser Asn Pro Ser 3890 3895 3900
- Ser Asn Val Ser Leu Gly Phe Glu Gly Cys Leu Asp Ala Val Val 3905 3910 3915 3920
- Asn Gly Glu Arg Leu Glu Leu Gly Arg Glu Lys Lys Met Glu Gly
  3925 3930 3935
- Arg Leu Glu Thr Trp Ala Leu Ser Gln Cys Cys Trp Pro Gly Thr Ala 3940 3945 3950
- Cys Ser Gln Ser Pro Cys Leu Asn Gly Gly Ser Cys Ser Pro Ala Leu 3955 3960 3965
- Gly Ser Gly Tyr Leu Cys Arg Cys Pro Pro Pro Phe Ser Gly Arg Asn 3970 3980
- Cys Glu Leu Gly Arg Glu Asn Cys Thr Ser Ala Pro Cys Gln Glu Gly 3985 3990 3995 4000

- Gly Thr Cys Val Ser Ser Pro Glu Gly Thr Ser Cys Asn Cys Pro His 4005 4010 4015
- Pro Tyr Thr Gly Asp Arg Cys Glu Met Glu Ala Arg Gly Cys Ser Gly 4020 4025 4030
- Gly His Cys Leu Ile Thr Pro Glu Ile Lys Arg Gly Asp Trp Gly Gln
  4035 4040 4045
- Gln Glu Phe Leu Val Ile Thr Val Ala Leu Pro Leu Val Ile Ile Ala 4050 4060
- Thr Val Gly Leu Leu Tyr Cys Arg Arg Arg Lys Ser His Lys Pro 4065 4070 4075 4080
- Val Thr Met Glu Asp Pro Asp Leu Leu Ala Arg Ser Ile Gly Val Asp 4085 4090 4095
- Thr Gln Ala Ser Pro Ala Ile Glu Leu Asp Pro Leu Asn Thr Ser Ser 4100 4105 4110
- Cys Asn Asn Leu Asn Gln Pro Glu Pro Ser Lys Thr Ser Val Pro Asn 4115 4120 4125
- Glu Leu Val Thr Phe Gly Pro Ser Ser Lys Gln Arg Pro Met Val Cys 4130 4135 4140
- Ser Val Pro Pro Arg Leu Pro Pro Ala Ala Val Ser Ser His Pro Gly 4145 4150 4155 4160
- His Glu Pro Ile Ile Lys Arg Thr Trp Ser Gly Glu Glu Leu Val Tyr 4165 4170 4175
- Pro Ser Gly Ala Ala Val Trp Pro Pro Thr Tyr Ser Arg Lys Lys His
  4180 4185 4190
- Trp Glu Tyr Pro His Pro Glu Thr Met Gln Gly Thr Leu Pro Pro Ser 4195 4200 4205
- Pro Arg Arg His Val Gly Pro Ala Val Met Pro Asp Pro Thr Gly Leu 4210 4215 4220
- Tyr Gly Gly Phe Pro Phe Pro Leu Glu Leu Glu Asn Lys Arg Ala Pro 4225 4230 4235 4240
- Leu Pro Pro Arg Tyr Ser Asn Gln Asn Leu Glu Asp Leu Met Pro Pro
  4245 4250 4255
- Arg Pro Pro Ser Pro Arg Glu His Leu Leu Ala Pro Cys Leu Asn Glu 4260 4265 4270
- Tyr Thr Ala Ile Ser Tyr Tyr His Ser Gln Phe Arg Gln Gly Gly 4275 4280 4285
- Gly Pro Cys Leu Ala Glu Gly Gly Tyr Lys Gly Val Ser Met Arg Leu 4290 4295 4300

Ser Arg Ala Gly Pro Ser Tyr Ala Asp Cys Glu Val Asn Gly Gly Pro 4305 4310 4315 4320

Ala Thr Gly Arg Ser Gln Pro Arg Ala Pro Pro Asn Tyr Glu Gly Ser 4325 4330 4335

Asp Met Val Glu Ser Asp Tyr Gly Ser Cys Glu Glu Val Met Phe 4340 4345 4350

<210> 21

<211> 4590

<212> PRT

<213> Homo sapiens

<400> 21

Met Gly Arg His Leu Ala Leu Leu Leu Leu Leu Leu Leu Leu Phe Gln

1 5 10 15

His Phe Gly Asp Ser Asp Gly Ser Gln Arg Leu Glu Gln Thr Pro Leu 20 25 30

Gln Phe Thr His Leu Glu Tyr Asn Val Thr Val Gln Glu Asn Ser Ala 35 40 45

Ala Lys Thr Tyr Val Gly His Pro Val Lys Met Gly Val Tyr Ile Thr 50 55 60

His Pro Ala Trp Glu Val Arg Tyr Lys Ile Val Ser Gly Asp Ser Glu 65 70 75 80

Asn Leu Phe Lys Ala Glu Glu Tyr Ile Leu Gly Asp Phe Cys Phe Leu 85 90 95

Arg Ile Arg Thr Lys Gly Gly Asn Thr Ala Ile Leu Asn Arg Glu Val

Lys Asp His Tyr Thr Leu Ile Val Lys Ala Leu Glu Lys Asn Thr Asn 115 120 125

Val Glu Ala Arg Thr Lys Val Arg Val Gln Val Leu Asp Thr Asn Asp 130 135 140

Leu Arg Pro Leu Phe Ser Pro Thr Ser Tyr Ser Val Ser Leu Pro Glu 145 150 155 160

Asn Thr Ala Ile Arg Thr Ser Ile Ala Arg Val Ser Ala Thr Asp Ala 165 170 175

Asp Ile Gly Thr Asn Gly Glu Phe Tyr Tyr Ser Phe Lys Asp Arg Thr
180 185 190

Asp Met Phe Ala Ile His Pro Thr Ser Gly Val Ile Val Leu Thr Gly 195 200 205

Arg Leu Asp Tyr Leu Glu Thr Lys Leu Tyr Glu Met Glu Ile Leu Ala

	2.5	
210	215	220

Ala 225	Asp	Arg	Gly	Met	Lys 230	Leu	Tyr	Gly	Ser	Ser 235	Gly	Ile	Ser	Ser	Met 240
Ala	Lys	Leu	Thr	Val 245	His	Ile	Glu	Gln	Ala 250	Asn	Glu	Cys	Ala	Pro 255	Val
Ile	Thr	Ala	Val 260	Thr	Leu	Ser	Pro	Ser 265	Glu	Leu	Asp	Arg	Asp 270	Pro	Ala
Tyr	Ala	Ile 275	Val	Thr	Val	Asp	Asp 280	Cys	Asp	Gln	Gly	Ala 285	Asn	Gly	Asp
Ile	Ala 290	Ser	Leu	Ser	Ile	Val 295	Ala	Gly	Asp	Leu	Leu 300	Gln	Gln	Phe	Arg
Thr 305	Val	Arg	Ser	Phe	Pro 310	Gly	Ser	Lys	Glu	Tyr 315	Lys	Val	Lys	Ala	Ile 320
Gly	Asp	Ile	Asp	Trp 325	Asp	Ser	His	Pro	Phe 330	Gly	Tyr	Asn	Leu	Thr 335	Leu
Gln	Ala	Lys	Asp 340	Lys	Gly	Thr	Pro	Pro 345	Gln	Phe	Ser	Ser	Val 350	Lys	Val
Ile	His	Val 355	Thr	Ser	Pro	Gln	Phe 360	Lys	Ala	Gly	Pro	Val 365	Lys	Phe	Glu
Lys	Asp 370	Val	Tyr	Arg	Ala	Glu 375	Ile	Ser	Glu	Phe	Ala 380	Pro	Pro	Asn	Thr
Pro 385	Val	Val	Met	Val	Lys 390	Ala	Ile	Pro	Ala	Tyr 395	Ser	His	Leu	Arg	Tyr 400
Val	Phe	Lys	Arg	Thr 405	Pro	Gly	Lys	Ala	Lys 410	Phe	Ser	Leu	Asn	Tyr 415	Asn
Thr	Gly	Leu	Ile 420	Ser	Ile	Leu	Glu	Pro 425	Val	Lys	Arg	Gln	Gln 430	Ala	Ala
His	Phe	Glu 435	Leu	Glu	Val	Thr	Thr 440	Ser	Asp	Arg	Lys	Ala 445	Ser	Thr	Lys
Val	Leu 450	Val	Lys	Val	Leu	Gly 455	Ala	Asn	Ser	Asn	Pro 460	Pro	Glu	Phe	Thr
Gln 465	Thr	Ala	Tyr	Lys	Ala 470	Ala	Phe	Asp	Glu	Asn 475	Val	Pro	Ile	Gly	Thr 480
Thr	Ile	Met	Ser	Leu 485	Ser	Ala	Val	Asp	Pro 490	Asp	Glu	Gly	Glu	Asn 495	Gly
Tyr	Val	Thr	Tyr 500	Ser	Ile	Ala	Asn	Leu 505	Asn	His	Val	Pro	Phe 510	Ala	Ile
Asp	His	Phe	Thr	Gly	Ala	Val	Ser	Thr	Ser	Glu	Asn	Leu	Asp	Tyr	Glu

Leu Met Pro Arg Val Tyr Thr Leu Arg Ile Arg Ala Ser Asp Trp Gly 535 Leu Pro Tyr Arg Arg Glu Val Glu Val Leu Ala Thr Ile Thr Leu Asn 545 550 Asn Leu Asn Asp Asn Thr Pro Leu Phe Glu Lys Ile Asn Cys Glu Gly 570 Thr Ile Pro Arg Asp Leu Gly Val Gly Glu Gln Ile Thr Thr Val Ser 580 Ala Ile Asp Ala Asp Glu Leu Gln Leu Val Gln Tyr Gln Ile Glu Ala 600 Gly Asn Glu Leu Asp Leu Phe Ser Leu Asn Pro Asn Ser Gly Val Leu 615 Ser Leu Lys Arg Ser Leu Met Asp Gly Leu Gly Ala Lys Val Ser Phe 625 630 His Ser Leu Arg Ile Thr Ala Thr Asp Gly Glu Asn Phe Ala Thr Pro 650 Leu Tyr Ile Asn Ile Thr Val Ala Ala Ser His Lys Leu Val Asn Leu 660 Gln Cys Glu Glu Thr Gly Val Ala Lys Met Leu Ala Glu Lys Leu Leu Gln Ala Asn Lys Leu His Asn Gln Gly Glu Val Glu Asp Ile Phe Phe 695 Asp Ser His Ser Val Asn Ala His Ile Pro Gln Phe Arg Ser Thr Leu 705 Pro Thr Gly Ile Gln Val Lys Glu Asn Gln Pro Val Gly Ser Ser Val Ile Phe Met Asn Ser Thr Asp Leu Asp Thr Gly Phe Asn Gly Lys Leu 740 745 Val Tyr Ala Val Ser Gly Gly Asn Glu Asp Ser Cys Phe Met Ile Asp Met Glu Thr Gly Met Leu Lys Ile Leu Ser Pro Leu Asp Arg Glu Thr 775 Thr Asp Lys Tyr Thr Leu Asn Ile Thr Val Tyr Asp Leu Gly Ile Pro 785 800 Gln Lys Ala Ala Trp Arg Leu Leu His Val Val Val Asp Ala Asn

Asp Asn Pro Pro Glu Phe Leu Gln Glu Ser Tyr Phe Val Glu Val Ser

820	825	830

Glu Asp Lys 835	Glu Val	His Ser	Glu I 840	Ile Ile	Gln Val	Glu Ala 845	Thr	Asp
Lys Asp Leu 850	Gly Pro	Asn Gly 855	His V	/al Thr	Tyr Ser 860	Ile Leu	Thr	Asp
Thr Asp Thr 865	Phe Ser	Ile Asp 870	Ser V	al Thr	Gly Val 875	Val Asn	Ile	Ala 880
Arg Pro Leu	Asp Arg 885	Glu Leu	Gln H	His Glu 890	His Ser	Leu Lys	Ile 895	Glu
Ala Arg Asp	Gln Ala 900	Arg Glu		Pro Gln 905	Leu Phe	Ser Thr 910	Val	Val
Val Lys Val 915	Ser Leu	Glu Asp	Val A 920	Asn Asp	Asn Pro	Pro Thr 925	Phe	Ile
Pro Pro Asn 930	Tyr Arg	Val Lys 935	Val A	Arg Glu	Asp Leu 940	Pro Glu	Gly	Thr
Val Ile Met 945	Trp Leu	Glu Ala 950	His A	Asp Pro	Asp Leu 955	Gly Gln		Gly 960
Gln Val Arg	Tyr Ser 965	Leu Leu	Asp H	His Gly 970	Glu Gly	Asn Phe	Asp 975	Val
Asp Lys Leu	Ser Gly 980	Ala Val	_	lle Val	Gln Gln	Leu Asp 990	Phe	Glu
Lys Lys Gln 995	Val Tyr		Thr V 1000	/al Arg	_	Asp Lys 1005	Gly	Lys
Pro Val Ser 1010	Leu Ser	Ser Thr 1015	Cys T	Cyr Val	Glu Val 1020	Glu Val	Val .	Asp
Val Asn Glu 1025		His Pro 1030	Pro V		Ser Ser .035	Phe Val		Lys 040
Gly Thr Val	Lys Glu 1045	Asp Ala	Pro V	al Gly 1050	Ser Leu		Thr 1	Val
Ser Ala His	Asp Glu 1060	Asp Ala	-	arg Asp 165	Gly Glu	Ile Arg 1070	Tyr	Ser
Ile Arg Asp 1075	Gly Ser	=	Gly V 1080	al Phe	_	Gly Glu .085	Glu '	Thr
Gly Val Ile 1090	Glu Thr	Ser Asp 1095	Arg L	eu Asp	Arg Glu 1100	Ser Thr	Ser 1	His
Tyr Trp Leu 1105		Phe Ala 1110	Thr A		Gly Val 115	Val Pro		Ser 120

Ser Phe Ile Glu Ile Tyr Ile Glu Val Glu Asp Val Asn Asp Asn Ala

- Pro Gln Thr Ser Glu Pro Val Tyr Tyr Pro Glu Ile Met Glu Asn Ser 1140 1145 1150
- Pro Lys Asp Val Ser Val Val Gln Ile Glu Ala Phe Asp Pro Asp Ser 1155 1160 1165
- Ser Ser Asn Asp Lys Leu Met Tyr Lys Ile Thr Ser Gly Asn Pro Gln 1170 1175 1180
- Gly Phe Phe Ser Ile His Pro Lys Thr Gly Leu Ile Thr Thr Ser 1185 1190 1195 1200
- Arg Lys Leu Asp Arg Glu Gln Gln Asp Glu His Ile Leu Glu Val Thr 1205 1210 1215
- Val Thr Asp Asn Gly Ser Pro Pro Lys Ser Thr Ile Ala Arg Val Ile 1220 1225 1230
- Val Lys Ile Leu Asp Glu Asn Asp Asn Lys Pro Gln Phe Leu Gln Lys 1235 1240 1245
- Phe Tyr Lys Ile Arg Leu Pro Glu Arg Glu Lys Pro Asp Arg Glu Arg 1250 1255 1260
- Asn Ala Arg Arg Glu Pro Leu Tyr Arg Val Ile Ala Thr Asp Lys Asp 1265 1270 1275 1280
- Glu Gly Pro Asn Ala Glu Ile Ser Tyr Ser Ile Glu Asp Gly Asn Glu 1285 1290 1295
- His Gly Lys Phe Phe Ile Glu Pro Lys Thr Gly Val Val Ser Ser Lys
  1300 1305 1310
- Arg Phe Ser Ala Ala Gly Glu Tyr Asp Ile Leu Ser Ile Lys Ala Val 1315 1320 1325
- Asp Asn Gly Arg Pro Gln Lys Ser Ser Thr Thr Arg Leu His Ile Glu
  1330 1340
- Trp Ile Ser Lys Pro Lys Gln Ser Leu Glu Pro Ile Ser Phe Glu Glu 1345 1350 1355 1360
- Ser Phe Phe Thr Phe Thr Val Met Glu Ser Asp Pro Val Ala His Met 1365 1370 1375
- Ile Gly Val Ile Ser Val Glu Pro Pro Gly Ile Pro Leu Trp Phe Asp 1380 1385 1390
- Ile Thr Gly Gly Asn Tyr Asp Ser His Phe Asp Val Asp Lys Gly Thr
  1395 1400 1405
- Gly Thr Ile Ile Val Ala Lys Pro Leu Asp Ala Glu Gln Lys Ser Asn 1410 1415 1420
- Tyr Asn Leu Thr Val Glu Ala Thr Asp Gly Thr Thr Thr Ile Leu Thr

- Gln Val Phe Ile Lys Val Ile Asp Thr Asn Asp His Arg Pro Gln Phe 1445 1450 1455
- Ser Thr Ser Lys Tyr Glu Val Val Ile Pro Glu Asp Thr Ala Pro Glu 1460 1465 1470
- Thr Glu Ile Leu Gln Ile Ser Ala Val Asp Gln Asp Glu Lys Asn Lys 1475 1480 1485
- Leu Ile Tyr Thr Leu Gln Ser Ser Arg Asp Pro Leu Ser Leu Lys Lys 1490 1495 1500
- Phe Arg Leu Asp Pro Ala Thr Gly Ser Leu Tyr Thr Ser Glu Lys Leu 1505 1510 1515 1520
- Asp His Glu Ala Val Ser Pro Ala His Leu Thr Val Met Val Arg Asp 1525 1530 1535
- Gln Asp Val Pro Val Lys Arg Asn Phe Ala Arg Ile Val Val Asn Val 1540 1545 1550
- Ser Asp Thr Asn Asp His Ala Pro Trp Phe Thr Ala Ser Ser Tyr Lys 1555 1560 1565
- Gly Arg Val Tyr Glu Ser Ala Ala Val Gly Ser Val Val Leu Gln Val 1570 1580
- Thr Ala Leu Asp Lys Asp Lys Gly Lys Asn Ala Glu Val Leu Tyr Ser 1585 1590 1595 1600
- Ile Glu Ser Gly Asn Ile Gly Asn Ile Gly Asn Ser Phe Met Ile Asp 1605 1610 1615
- Pro Val Leu Gly Ser Ile Lys Thr Ala Lys Glu Leu Asp Arg Ser Asn 1620 1625 1630
- Gln Ala Glu Tyr Asp Leu Met Val Lys Ala Thr Asp Lys Gly Ser Pro 1635 1640 1645
- Pro Met Ser Glu Ile Thr Ser Val Arg Ile Phe Val Thr Ile Ala Asp 1650 1655 1660
- Asn Ala Ser Pro Lys Phe Thr Ser Lys Glu Tyr Ser Val Glu Leu Ser 1665 1670 1675 1680
- Glu Thr Val Ser Ile Gly Ser Phe Val Gly Met Val Thr Ala His Ser 1685 1690 1695
- Gln Ser Ser Val Val Tyr Glu Ile Lys Asp Gly Asn Thr Gly Asp Ala 1700 1705 1710
- Phe Asp Ile Asn Pro His Ser Gly Thr Ile Ile Thr Gln Lys Ala Leu 1715 1720 1725
- Asp Phe Glu Thr Leu Pro Ile Tyr Thr Leu Ile Ile Gln Gly Thr Asn

1730

- Met Ala Gly Leu Ser Thr Asn Thr Thr Val Leu Val His Leu Gln Asp 1745 1750 1755 1760
- Glu Asn Asp Asn Ala Pro Val Phe Met Gln Ala Glu Tyr Thr Gly Leu 1765 1770 1775
- Ile Ser Glu Ser Ala Ser Ile Asn Ser Val Val Leu Thr Asp Arg Asn 1780 1785 1790
- Val Pro Leu Val Ile Arg Ala Ala Asp Ala Asp Lys Asp Ser Asn Ala 1795 1800 1805
- Leu Leu Val Tyr His Ile Val Glu Pro Ser Val His Thr Tyr Phe Ala 1810 1815 1820
- Ile Asp Ser Ser Thr Gly Ala Ile His Thr Val Leu Ser Leu Asp Tyr 1825 1830 1835 1840
- Glu Glu Thr Ser Ile Phe His Phe Thr Val Gln Val His Asp Met Gly 1845 1850 1855
- Thr Pro Arg Leu Phe Ala Glu Tyr Ala Ala Asn Val Thr Val His Val
  1860 1865 1870
- Ile Asp Ile Asn Asp Cys Pro Pro Val Phe Ala Lys Pro Leu Tyr Glu 1875 1880 1885
- Ala Ser Leu Leu Pro Thr Tyr Lys Gly Val Lys Val Ile Thr Val 1890 1895 1900
- Asn Ala Thr Asp Ala Asp Ser Ser Ala Phe Ser Gln Leu Ile Tyr Ser 1905 1910 1915 1920
- Ile Thr Glu Gly Asn Ile Gly Glu Lys Phe Ser Met Asp Tyr Lys Thr 1925 1930 1935
- Gly Ala Leu Thr Val Gln Asn Thr Thr Gln Leu Arg Ser Arg Tyr Glu 1940 1945 1950
- Leu Thr Val Arg Ala Ser Asp Gly Arg Phe Ala Gly Leu Thr Ser Val 1955 1960 1965
- Lys Ile Asn Val Lys Glu Ser Lys Glu Ser His Leu Lys Phe Thr Gln 1970 1975 1980
- Asp Val Tyr Ser Ala Val Val Lys Glu Asn Ser Thr Glu Ala Glu Thr 1985 1990 1995 2000
- Leu Ala Val Ile Thr Ala Ile Gly Ser Pro Ile Asn Glu Pro Leu Phe 2005 2010 2015
- Tyr His Ile Leu Asn Pro Asp Arg Arg Phe Lys Ile Ser Arg Thr Ser 2020 2025 2030
- Gly Val Leu Ser Thr Thr Gly Thr Pro Phe Asp Arg Glu Gln Glu

- Ala Phe Asp Val Val Glu Val Ile Glu Glu His Lys Pro Ser Ala 2050 2055 2060
- Val Ala His Val Val Lys Val Ile Val Glu Asp Gln Asn Asp Asn 2065 2070 2075 2080
- Ala Pro Val Phe Val Asn Leu Pro Tyr Tyr Ala Val Val Lys Val Asp 2085 2090 2095
- Thr Glu Val Gly His Val Ile Arg Tyr Val Thr Ala Val Asp Arg Asp 2100 2105 2110
- Ser Gly Arg Asn Gly Glu Val His Tyr Tyr Leu Lys Glu His His Glu 2115 2120 2125
- His Phe Gln Ile Gly Pro Leu Gly Glu Ile Ser Leu Lys Lys Gln Phe 2130 2135 2140
- Glu Leu Asp Thr Leu Asn Lys Glu Tyr Leu Val Thr Val Val Ala Lys 2145 2150 2155 2160
- Asp Gly Gly Asn Pro Ala Phe Ser Ala Glu Val Ile Val Pro Ile Thr 2165 2170 2175
- Val Met Asn Lys Ala Met Pro Val Phe Glu Lys Pro Phe Tyr Ser Ala 2180 2185 2190
- Glu Ile Ala Glu Ser Ile Gln Val His Ser Pro Val Val His Val Gln 2195 2200 2205
- Ala Asn Ser Pro Glu Gly Leu Lys Val Phe Tyr Ser Ile Thr Asp Gly 2210 2215 2220
- Asp Pro Phe Ser Gln Phe Thr Ile Asn Phe Asn Thr Gly Val Ile Asn 2225 2230 2235 2240
- Val Ile Ala Pro Leu Asp Phe Glu Ala His Pro Ala Tyr Lys Leu Ser 2245 2250 2255
- Ile Arg Ala Thr Asp Ser Leu Thr Gly Ala His Ala Glu Val Phe Val 2260 2265 2270
- Asp Ile Ile Val Asp Asp Ile Asn Asp Asn Pro Pro Val Phe Ala Gln 2275 2280 2285
- Gln Ser Tyr Ala Val Thr Leu Ser Glu Ala Ser Val Ile Gly Thr Ser 2290 2295 2300
- Val Val Gln Val Arg Ala Thr Asp Ser Asp Ser Glu Pro Asn Arg Gly 2305 2310 2315 2320
- Ile Ser Tyr Gln Met Phe Gly Asn His Ser Lys Ser His Asp His Phe
  2325 2330 2335
- His Val Asp Ser Ser Thr Gly Leu Ile Ser Leu Leu Arg Thr Leu Asp

- Tyr Glu Gln Ser Arg Gln His Thr Ile Phe Val Arg Ala Val Asp Gly 2355 2360 2365
- Gly Met Pro Thr Leu Ser Ser Asp Val Ile Val Thr Val Asp Val Thr 2370 2375 2380
- Asp Leu Asn Gly Asn Pro Pro Leu Phe Glu Gln Gln Ile Tyr Glu Ala 2385 2390 2395 2400
- Arg Ile Ser Glu His Ala Pro His Gly His Phe Val Thr Cys Val Lys 2405 2410 2415
- Ala Tyr Asp Ala Asp Ser Ser Asp Ile Asp Lys Leu Gln Tyr Ser Ile 2420 2425 2430
- Leu Ser Gly Asn Asp His Lys His Phe Val Ile Asp Ser Ala Thr Gly 2435 2440 2445
- Ile Ile Thr Leu Ser Asn Leu His Arg His Ala Leu Lys Pro Phe Tyr 2450 2455 2460
- Ser Leu Asn Leu Ser Val Ser Asp Gly Val Phe Arg Ser Ser Thr Gln 2465 2470 2475 2480
- Val His Val Thr Val Ile Gly Gly Asn Leu His Ser Pro Ala Phe Leu 2485 2490 2495
- Gln Asn Glu Tyr Glu Val Glu Leu Ala Glu Asn Ala Pro Leu His Thr 2500 2505 2510
- Leu Val Met Glu Val Lys Thr Thr Asp Gly Asp Ser Gly Ile Tyr Gly 2515 2520 2525
- His Val Thr Tyr His Ile Val Asn Asp Phe Ala Lys Asp Arg Phe Tyr 2530 2535 2540
- Ile Asn Glu Arg Gly Gln Ile Phe Thr Leu Glu Lys Leu Asp Arg Glu 2545 2550 2555 2560
- Thr Pro Ala Glu Lys Val Ile Ser Val Arg Leu Met Ala Lys Asp Ala 2565 2570 2575
- Gly Gly Lys Val Ala Phe Cys Thr Val Asn Val Ile Leu Thr Asp Asp 2580 2590
- Asn Asp Asn Ala Pro Gln Phe Arg Ala Thr Lys Tyr Glu Val Asn Ile 2595 2600 2605
- Gly Ser Ser Ala Ala Lys Gly Thr Ser Val Val Lys Ser Ala Ser Asp 2610 2615 2620
- Ala Asp Glu Gly Ser Asn Ala Asp Ile Thr Tyr Ala Ile Glu Ala Asp 2625 2630 2635 2640
- Ser Glu Ser Val Lys Glu Asn Leu Glu Ile Asn Lys Leu Ser Gly Val

- Ile Thr Thr Lys Glu Ser Leu Ile Gly Leu Glu Asn Glu Phe Phe Thr 2660 2665 2670
- Phe Phe Val Arg Ala Val Asp Asn Gly Ser Pro Ser Lys Glu Ser Val 2675 2680 2685
- Val Leu Val Tyr Val Lys Ile Leu Pro Pro Glu Met Gln Leu Pro Lys 2690 2695 2700
- Phe Ser Glu Pro Phe Tyr Thr Phe Thr Val Ser Glu Asp Val Pro Val 2705 2710 2715 2720
- Gly Thr Glu Ile Asp Leu Ile Arg Ala Glu His Ser Gly Thr Val Leu 2725 2730 2735
- Tyr Ser Leu Val Lys Gly Asn Thr Pro Glu Ser Asn Arg Asp Glu Ser 2740 2745 2750
- Phe Val Ile Asp Arg Gln Ser Gly Arg Leu Lys Leu Glu Lys Ser Leu 2755 2760 2765
- Asp His Glu Thr Thr Lys Trp Tyr Gln Phe Ser Ile Leu Ala Arg Cys 2770 2775 2780
- Thr Gln Asp Asp His Glu Met Val Ala Ser Val Asp Val Ser Ile Gln 2785 2790 2795 2800
- Val Lys Asp Ala Asn Asp Asn Ser Pro Val Phe Glu Ser Ser Pro Tyr 2805 2810 2815
- Glu Ala Phe Ile Val Glu Asn Leu Pro Gly Gly Ser Arg Val Ile Gln 2820 2825 2830
- Ile Arg Ala Ser Asp Ala Asp Ser Gly Thr Asn Gly Gln Val Met Tyr 2835 2840 2845
- Ser Leu Asp Gln Ser Gln Ser Val Glu Val Ile Glu Ser Phe Ala Ile 2850 2855 2860
- Asn Met Glu Thr Gly Trp Ile Thr Thr Leu Lys Glu Leu Asp His Glu 2865 2870 2875 2880
- Lys Arg Asp Asn Tyr Gln Ile Lys Val Val Ala Ser Asp His Gly Glu 2885 2890 2895
- Lys Ile Gln Leu Ser Ser Thr Ala Ile Val Asp Val Thr Val Thr Asp 2900 2905 2910
- Val Asn Asp Ser Pro Pro Arg Phe Thr Ala Glu Ile Tyr Lys Gly Thr
  2915 2920 2925
- Val Ser Glu Asp Asp Pro Gln Gly Gly Val Ile Ala Ile Leu Ser Thr 2930 2935 2940
- Thr Asp Ala Asp Ser Glu Glu Ile Asn Arg Gln Val Thr Tyr Phe Ile

- Thr Gly Gly Asp Pro Leu Gly Gln Phe Ala Val Glu Thr Ile Gln Asn 2965 2970 2975
- Glu Trp Lys Val Tyr Val Lys Lys Pro Leu Asp Arg Glu Lys Arg Asp 2980 2985 2990
- Asn Tyr Leu Leu Thr Ile Thr Ala Thr Asp Gly Thr Phe Ser Ser Lys 2995 3000 3005
- Ala Ile Val Glu Val Lys Val Leu Asp Ala Asn Asp Asn Ser Pro Val 3010 3015 3020
- Cys Glu Lys Thr Leu Tyr Ser Asp Thr Ile Pro Glu Asp Val Leu Pro 3025 3030 3035 3040
- Gly Lys Leu Ile Met Gln Ile Ser Ala Thr Asp Ala Asp Ile Arg Ser 3045 3050 3055
- Asn Ala Glu Ile Thr Tyr Thr Leu Leu Gly Ser Gly Ala Glu Lys Phe 3060 3065 3070
- Lys Leu Asn Pro Asp Thr Gly Glu Leu Lys Thr Ser Thr Pro Leu Asp 3075 3080 3085
- Arg Glu Glu Gln Ala Val Tyr His Leu Leu Val Arg Ala Thr Asp Gly 3090 3095 3100
- Gly Gly Arg Phe Cys Gln Ala Ser Ile Val Val Thr Leu Glu Asp Val 3105 3110 3115 3120
- Asn Asp Asn Ala Pro Glu Phe Ser Ala Asp Pro Tyr Ala Ile Thr Val 3125 3130 3135
- Phe Glu Asn Thr Glu Pro Gly Thr Leu Leu Thr Arg Val Gln Ala Thr 3140 3145 3150
- Asp Ala Asp Ala Gly Leu Asn Arg Lys Ile Leu Tyr Ser Leu Ile Asp 3155 3160 3165
- Ser Ala Asp Gly Gln Phe Ser Ile Asn Glu Leu Ser Gly Ile Ile Gln 3170 3180
- Leu Glu Lys Pro Leu Asp Arg Glu Leu Gln Ala Val Tyr Thr Leu Ser 3185 3190 3195 3200
- Leu Lys Ala Val Asp Gln Gly Leu Pro Arg Arg Leu Thr Ala Thr Gly 3205 3210 3215
- Thr Val Ile Val Ser Val Leu Asp Ile Asn Asp Asn Pro Pro Val Phe 3220 3225 3230
- Glu Tyr Arg Glu Tyr Gly Ala Thr Val Ser Glu Asp Ile Leu Val Gly 3235 3240 3245
- Thr Glu Val Leu Gln Val Tyr Ala Ala Ser Arg Asp Ile Glu Ala Asn

- Ala Glu Ile Thr Tyr Ser Ile Ile Ser Gly Asn Glu His Gly Lys Phe 3265 3270 3275 3280
- Ser Ile Asp Ser Lys Thr Gly Ala Val Phe Ile Ile Glu Asn Leu Asp 3285 3290 3295
- Tyr Glu Ser Ser His Glu Tyr Tyr Leu Thr Val Glu Ala Thr Asp Gly 3300 3310
- Gly Thr Pro Ser Leu Ser Asp Val Ala Thr Val Asn Val Asn Val Thr
  3315 3320 3325
- Asp Ile Asn Asp Asn Thr Pro Val Phe Ser Gln Asp Thr Tyr Thr Thr 3330 3335 3340
- Val Ile Ser Glu Asp Ala Val Leu Glu Gln Ser Val Ile Thr Val Met 3345 3350 3355 3360
- Ala Asp Asp Ala Asp Gly Pro Ser Asn Ser His Ile His Tyr Ser Ile 3365 3370 3375
- Ile Asp Gly Asn Gln Gly Ser Ser Phe Thr Ile Asp Pro Val Arg Gly 3380 3385 3390
- Glu Val Lys Val Thr Lys Leu Leu Asp Arg Glu Thr Ile Ser Gly Tyr 3395 3400 3405
- Thr Leu Thr Val Gln Ala Ser Asp Asn Gly Ser Pro Pro Arg Val Asn 3410 3415 3420
- Thr Thr Thr Val Asn Ile Asp Val Ser Asp Val Asn Asp Asn Ala Pro 3425 3430 3435 3440
- Val Phe Ser Arg Gly Asn Tyr Ser Val Ile Ile Gln Glu Asn Lys Pro 3445 3450 3455
- Val Gly Phe Ser Val Leu Gln Leu Val Val Thr Asp Glu Asp Ser Ser 3460 3465 3470
- His Asn Gly Pro Pro Phe Phe Phe Thr Ile Val Thr Gly Asn Asp Glu 3475 3480 3485
- Lys Ala Phe Glu Val Asn Pro Gln Gly Val Leu Leu Thr Ser Ser Ala 3490 3495 3500
- Ile Lys Arg Lys Glu Lys Asp His Tyr Leu Leu Gln Val Lys Val Ala 3505 3510 3515 3520
- Asp Asn Gly Lys Pro Gln Leu Ser Ser Leu Thr Tyr Ile Asp Ile Arg 3525 3530 3535
- Val Ile Glu Glu Ser Ile Tyr Pro Pro Ala Ile Leu Pro Leu Glu Ile 3540 3545 3550
- Phe Ile Thr Ser Ser Gly Glu Glu Tyr Ser Gly Gly Val Ile Gly Lys

- Ile His Ala Thr Asp Gln Asp Val Tyr Asp Thr Leu Thr Tyr Ser Leu 3570 3580
- Asp Pro Gln Met Asp Asn Leu Phe Ser Val Ser Ser Thr Gly Gly Lys 3585 3590 3595 3600
- Leu Ile Ala His Lys Lys Leu Asp Ile Gly Gln Tyr Leu Leu Asn Val 3605 3610 3615
- Ser Val Thr Asp Gly Lys Phe Thr Thr Val Ala Asp Ile Thr Val His 3620 3625 3630
- Ile Arg Gln Val Thr Gln Glu Met Leu Asn His Thr Ile Ala Ile Arg 3635 3640 3645
- Phe Ala Asn Leu Thr Pro Glu Glu Phe Val Gly Asp Tyr Trp Arg Asn 3650 3655 3660
- Phe Gln Arg Ala Leu Arg Asn Ile Leu Gly Val Arg Arg Asn Asp Ile 3665 3670 3675 3680
- Gln Ile Val Ser Leu Gln Ser Ser Glu Pro His Pro His Leu Asp Val 3685 3690 3695
- Leu Leu Phe Val Glu Lys Pro Gly Ser Ala Gln Ile Ser Thr Lys Gln 3700 3705 3710
- Leu Leu His Lys Ile Asn Ser Ser Val Thr Asp Ile Glu Glu Ile Ile 3715 3720 3725
- Gly Val Arg Ile Leu Asn Val Phe Gln Lys Leu Cys Ala Gly Leu Asp 3730 3740
- Cys Pro Trp Lys Phe Cys Asp Glu Lys Val Ser Val Asp Glu Ser Val 3745 3750 3755 3760
- Met Ser Thr His Ser Thr Ala Arg Leu Ser Phe Val Thr Pro Arg His 3765 3770 3775
- His Arg Ala Ala Val Cys Leu Cys Lys Glu Gly Arg Cys Pro Pro Val 3780 3785 3790
- His His Gly Cys Glu Asp Asp Pro Cys Pro Glu Gly Ser Glu Cys Val 3795 3800 3805
- Ser Asp Pro Trp Glu Glu Lys His Thr Cys Val Cys Pro Ser Gly Arg 3810 3815 3820
- Phe Gly Gln Cys Pro Gly Ser Ser Ser Met Thr Leu Thr Gly Asn Ser 3825 3830 3835 3840
- Tyr Val Lys Tyr Arg Leu Thr Glu Asn Glu Asn Lys Leu Glu Met Lys 3845 3850 3855
- Leu Thr Met Arg Leu Arg Thr Tyr Ser Thr His Ala Val Val Met Tyr

- Ala Arg Gly Thr Asp Tyr Ser Ile Leu Glu Ile His His Gly Arg Leu 3875 3880 3885
- Gln Tyr Lys Phe Asp Cys Gly Ser Gly Pro Gly Ile Val Ser Val Gln 3890 3895 3900
- Ser Ile Gln Val Asn Asp Gly Gln Trp His Ala Val Ala Leu Glu Val 3905 3910 3915 3920
- Asn Gly Asn Tyr Ala Arg Leu Val Leu Asp Gln Val His Thr Ala Ser 3925 3930 3935
- Gly Thr Ala Pro Gly Thr Leu Lys Thr Leu Asn Leu Asp Asn Tyr Val 3940 3945 3950
- Phe Phe Gly Gly His Ile Arg Gln Gln Gly Thr Arg His Gly Arg Ser 3955 3960 3965
- Pro Gln Val Gly Asn Gly Phe Arg Gly Cys Met Asp Ser Ile Tyr Leu 3970 3980
- Asn Gly Gln Glu Leu Pro Leu Asn Ser Lys Pro Arg Ser Tyr Ala His 3985 3990 3995 4000
- Ile Glu Glu Ser Val Asp Val Ser Pro Gly Cys Phe Leu Thr Ala Thr 4005 4010 4015
- Glu Asp Cys Ala Ser Asn Pro Cys Gln Asn Gly Gly Val Cys Asn Pro 4020 4025 4030
- Ser Pro Ala Gly Gly Tyr Tyr Cys Lys Cys Ser Ala Leu Tyr Ile Gly 4035 4040 4045
- Thr His Cys Glu Ile Ser Val Asn Pro Cys Ser Ser Asn Pro Cys Leu 4050 4055 4060
- Tyr Gly Gly Thr Cys Val Val Asp Asn Gly Gly Phe Val Cys Gln Cys 4065 4070 4075 4080
- Arg Gly Leu Tyr Thr Gly Gln Arg Cys Gln Leu Ser Pro Tyr Cys Lys 4085 4090 4095
- Asp Glu Pro Cys Lys Asn Gly Gly Thr Cys Phe Asp Ser Leu Asp Gly 4100 4105 4110
- Ala Val Cys Gln Cys Asp Ser Gly Phe Arg Gly Glu Arg Cys Gln Ser 4115 4120 4125
- Asp Ile Asp Glu Cys Ser Gly Asn Pro Cys Leu His Gly Ala Leu Cys 4130 4140
- Glu Asn Thr His Gly Ser Tyr His Cys Asn Cys Ser His Glu Tyr Arg 4145 4150 4155 4160
- Gly Arg His Cys Glu Asp Ala Ala Pro Asn Gln Tyr Val Ser Thr Pro

- Trp Asn Ile Gly Leu Ala Glu Gly Ile Gly Ile Val Val Phe Val Ala 4180 4185 4190
- Gly Ile Phe Leu Leu Val Val Val Phe Val Leu Cys Arg Lys Met Ile 4195 4200 4205
- Ser Arg Lys Lys His Gln Ala Glu Pro Lys Asp Lys His Leu Gly 4210 4215 4220
- Pro Ala Thr Ala Phe Leu Gln Arg Pro Tyr Phe Asp Ser Lys Leu Asn 4225 4230 4235 4240
- Lys Asn Ile Tyr Ser Asp Ile Pro Pro Gln Val Pro Val Arg Pro Ile 4245 4250 4255
- Ser Tyr Thr Pro Ser Ile Pro Ser Asp Ser Arg Asn Asn Leu Asp Arg 4260 4265 4270
- Asn Ser Phe Glu Gly Ser Ala Ile Pro Glu His Pro Glu Phe Ser Thr 4275 4280 4285
- Phe Asn Pro Glu Ser Val His Gly His Arg Lys Ala Val Ala Val Cys 4290 4295 4300
- Ser Val Ala Pro Asn Leu Pro Pro Pro Pro Pro Ser Asn Ser Pro Ser 4305 4310 4315 4320
- Asp Ser Asp Ser Ile Gln Lys Pro Ser Trp Asp Phe Asp Tyr Asp Thr 4325 4330 4335
- Lys Val Val Asp Leu Asp Pro Cys Leu Ser Lys Lys Pro Leu Glu Glu 4340 4345 4350
- Lys Pro Ser Gln Pro Tyr Ser Ala Arg Glu Ser Leu Ser Glu Val Gln
  4355 4360 4365
- Ser Leu Ser Ser Phe Gln Ser Glu Ser Cys Asp Asp Asn Gly Tyr His 4370 4380
- Trp Asp Thr Ser Asp Trp Met Pro Ser Val Pro Leu Pro Asp Ile Gln 4385 4390 4395 4400
- Glu Phe Pro Asn Tyr Glu Val Ile Asp Glu Gln Thr Pro Leu Tyr Ser 4405 4410 4415
- Ala Asp Pro Asn Ala Ile Asp Thr Asp Tyr Tyr Pro Gly Gly Tyr Asp 4420 4425 4430
- Ile Glu Ser Asp Phe Pro Pro Pro Pro Glu Asp Phe Pro Ala Ala Asp 4435 4440 4445
- Glu Leu Pro Pro Leu Pro Pro Glu Phe Ser Asn Gln Phe Glu Ser Ile 4450 4455 4460
- His Pro Pro Arg Asp Met Pro Ala Ala Gly Ser Leu Gly Ser Ser Ser

Arg Asn Arg Gln Arg Phe Asn Leu Asn Gln Tyr Leu Pro Asn Phe Tyr 4485 4490 4495

Pro Leu Asp Met Ser Glu Pro Gln Thr Lys Gly Thr Gly Glu Asn Ser 4500 4505 4510

Thr Cys Arg Glu Pro His Ala Pro Tyr Pro Pro Gly Tyr Gln Arg His 4515 4520 4525

Phe Glu Ala Pro Ala Val Glu Ser Met Pro Met Ser Val Tyr Ala Ser 4530 4540

Thr Ala Ser Cys Ser Asp Val Ser Ala Cys Cys Glu Val Glu Ser Glu 4545 4550 4555 4560

Val Met Met Ser Asp Tyr Glu Ser Gly Asp Asp Gly His Phe Glu Glu 4565 4570 4575

Val Thr Ile Pro Pro Leu Asp Ser Gln Gln His Thr Glu Val 4580 4585 4590

<210> 22

4465

<211> 2991

<212> PRT

<213> Homo sapiens

<400> 22

Met Glu Thr Asp Pro Val Asn His Met Val Gly Val Ile Ser Val Glu

1 5 10 15

Gly Arg Pro Gly Leu Phe Trp Phe Asn Ile Ser Gly Gly Asp Lys Asp 20 25 30

Met Asp Phe Asp Ile Glu Lys Thr Thr Gly Ser Ile Val Ile Ala Arg 35 40 45

Pro Leu Asp Thr Arg Arg Arg Ser Asn Tyr Asn Leu Thr Val Glu Val
50 55 60

Thr Asp Gly Ser Arg Thr Ile Ala Thr Gln Val His Ile Phe Met Ile
65 70 75 80

Ala Asn Ile Asn His His Arg Pro Gln Phe Leu Glu Thr Arg Tyr Glu 85 90 95

Val Arg Val Pro Gln Asp Thr Val Pro Gly Val Glu Leu Leu Arg Val
100 105 110

Gln Ala Ile Asp Gln Asp Lys Gly Lys Ser Leu Ile Tyr Thr Ile His
115 120 125

Gly Ser Gln Asp Pro Gly Ser Ala Ser Leu Phe Gln Leu Asp Pro Ser 130 135 140

Ser Gly Val Leu Val Thr Val Gly Lys Leu Asp Leu Gly Ser Gly Pro 145 Ser Gln His Thr Leu Thr Val Met Val Arg Asp Gln Glu Ile Pro Ile 170 Lys Arg Asn Phe Val Trp Val Thr Ile His Val Glu Asp Gly Asn Leu His Pro Pro Arg Phe Thr Gln Leu His Tyr Glu Ala Ser Val Pro Asp 200 Thr Ile Ala Pro Gly Thr Glu Leu Leu Gln Val Arg Ala Met Asp Ala 215 Asp Arg Gly Val Asn Ala Glu Val His Tyr Ser Leu Leu Lys Gly Asn 230 235 Ser Glu Gly Phe Phe Asn Ile Asn Ala Leu Leu Gly Ile Ile Thr Leu Ala Gln Lys Leu Asp Gln Ala Asn His Ala Pro His Thr Leu Thr Val Lys Ala Glu Asp Gln Gly Ser Pro Gln Trp His Asp Leu Ala Thr Val 275 280 285 Ile Ile His Val Tyr Pro Ser Asp Arg Ser Ala Pro Ile Phe Ser Lys Ser Glu Tyr Phe Val Glu Ile Pro Glu Ser Ile Pro Val Gly Ser Pro 310 315 Ile Leu Leu Val Ser Ala Met Ser Pro Ser Glu Val Thr Tyr Glu Leu 325 Arg Glu Gly Asn Lys Asp Gly Val Phe Ser Met Asn Ser Tyr Ser Gly 345 Leu Ile Ser Thr Gln Lys Lys Leu Asp His Glu Lys Ile Ser Ser Tyr 360 Gln Leu Lys Ile Arg Gly Ser Asn Met Ala Gly Ala Phe Thr Asp Val Met Val Val Asp Ile Ile Asp Glu Asn Asp Asn Ala Pro Met Phe 390 Leu Lys Ser Thr Phe Val Gly Gln Ile Ser Glu Ala Ala Pro Leu Tyr 405 Ser Met Ile Met Asp Lys Asn Asn Pro Phe Val Ile His Ala Ser 425 Asp Ser Asp Lys Glu Ala Asn Ser Leu Leu Val Tyr Lys Ile Leu Glu 435 440 445

Pro Glu Ala Leu Lys Phe Phe Lys Ile Asp Pro Ser Met Gly Thr Leu Thr Ile Val Ser Glu Met Asp Tyr Glu Ser Met Pro Ser Phe Gln Phe Cys Val Tyr Val His Asp Gln Gly Ser Pro Val Leu Phe Ala Pro Arg 490 Pro Ala Gln Val Ile Ile His Val Arg Asp Val Asn Asp Ser Pro Pro 500 505 Arg Phe Ser Glu Gln Ile Tyr Glu Val Ala Ile Val Gly Pro Ile His 520 Pro Gly Met Glu Leu Leu Met Val Arg Ala Ser Asp Glu Asp Ser Glu 535 Val Asn Tyr Ser Ile Lys Thr Gly Asn Ala Asp Glu Ala Val Thr Ile His Pro Val Thr Gly Ser Ile Ser Val Leu Asn Pro Ala Phe Leu Gly 565 Leu Ser Arg Lys Leu Thr Ile Arg Ala Ser Asp Gly Leu Tyr Gln Asp 580 585 Thr Ala Leu Val Lys Ile Ser Leu Thr Gln Val Leu Asp Lys Ser Leu Gln Phe Asp Gln Asp Val Tyr Trp Ala Ala Val Lys Glu Asn Leu Gln Asp Arg Lys Ala Leu Val Ile Leu Gly Ala Gln Gly Asn His Leu Asn 630 Asp Thr Leu Ser Tyr Phe Leu Leu Asn Gly Thr Asp Met Phe His Met 650 Val Gln Ser Ala Gly Val Leu Gln Thr Arg Gly Val Ala Phe Asp Arg Glu Gln Gln Asp Thr His Glu Leu Ala Val Glu Val Arg Asp Asn Arg Thr Pro Gln Arg Val Ala Gln Gly Leu Val Arg Val Ser Ile Glu Asp Val Asn Asp Asn Pro Pro Lys Phe Lys His Leu Pro Tyr Tyr Thr Ile 710 Ile Gln Asp Gly Thr Glu Pro Gly Asp Val Leu Phe Gln Val Ser Ala 725 730 Thr Asp Glu Asp Leu Gly Thr Asn Gly Ala Val Thr Tyr Glu Phe Ala 745

Glu Asp Tyr Thr Tyr Phe Arg Ile Asp Pro Tyr Leu Gly Asp Ile Ser 755 Leu Lys Lys Pro Phe Asp Tyr Gln Ala Leu Asn Lys Tyr His Leu Lys Val Ile Ala Arg Asp Gly Gly Thr Pro Ser Leu Gln Ser Glu Glu Glu 790 Val Leu Val Thr Val Arg Asn Lys Ser Asn Pro Leu Phe Gln Ser Pro 810 Tyr Tyr Lys Val Arg Val Pro Glu Asn Ile Thr Leu Tyr Thr Pro Ile 825 Leu His Thr Gln Ala Arg Ser Pro Glu Gly Leu Arg Leu Ile Tyr Asn 840 Ile Val Glu Glu Pro Leu Met Leu Phe Thr Thr Asp Phe Lys Thr 855 Gly Val Leu Thr Val Thr Gly Pro Leu Asp Tyr Glu Ser Lys Thr Lys 870 His Val Phe Thr Val Arg Ala Thr Asp Thr Ala Leu Gly Ser Phe Ser 885 890 Glu Ala Thr Val Glu Val Leu Val Glu Asp Val Asn Asp Asn Pro Pro Thr Phe Ser Gln Leu Val Tyr Thr Thr Ser Ile Ser Glu Gly Leu Pro 920 Ala Gln Thr Pro Val Ile Gln Leu Leu Ala Ser Asp Gln Asp Ser Gly 930 935 Arg Asn Arg Asp Val Ser Tyr Gln Ile Val Glu Asp Gly Ser Asp Val 950 955 Ser Lys Phe Phe Gln Ile Asn Gly Ser Thr Gly Glu Met Ser Thr Val 970 Gln Glu Leu Asp Tyr Glu Ala Gln Gln His Phe His Val Lys Val Arg Ala Met Asp Lys Gly Asp Pro Pro Leu Thr Gly Glu Thr Leu Val Val 1000 Val Asn Val Ser Asp Ile Asn Asp Asn Pro Pro Glu Phe Arg Gln Pro 1010 1015

Gln Tyr Glu Ala Asn Val Ser Glu Leu Ala Thr Cys Gly His Leu Val

Leu Lys Val Gln Ala Ile Asp Pro Asp Ser Arg Asp Thr Ser Arg Leu

1030

1045

1050

1035

- Glu Tyr Leu Ile Leu Ser Gly Asn Gln Asp Arg His Phe Phe Ile Asn 1060 1065 1070
- Ser Ser Ser Gly Ile Ile Ser Met Phe Asn Leu Cys Lys Lys His Leu 1075 1080 1085
- Asp Ser Ser Tyr Asn Leu Arg Val Gly Ala Ser Asp Gly Val Phe Arg 1090 1095 1100
- Ala Thr Val Pro Val Tyr Ile Asn Thr Thr Asn Ala Asn Lys Tyr Ser 1105 1110 1115 1120
- Pro Glu Phe Gln Gln His Leu Tyr Glu Ala Glu Leu Ala Glu Asn Ala 1125 1130 1135
- Met Val Gly Thr Lys Val Ile Asp Leu Leu Ala Ile Asp Lys Asp Ser 1140 1145 1150
- Gly Pro Tyr Gly Thr Ile Asp Tyr Thr Ile Ile Asn Lys Leu Ala Ser 1155 1160 1165
- Glu Lys Phe Ser Ile Asn Pro Asn Gly Gln Ile Ala Thr Leu Gln Lys 1170 1175 1180
- Leu Asp Arg Glu Asn Ser Thr Glu Arg Val Ile Ala Ile Lys Val Met 1185 1190 1195 1200
- Ala Arg Asp Gly Gly Arg Val Ala Phe Cys Thr Val Lys Ile Ile 1205 1210 1215
- Leu Thr Asp Glu Asn Asp Asn Pro Pro Gln Phe Lys Ala Ser Glu Tyr
  1220 1225 1230
- Thr Val Ser Ile Gln Ser Asn Val Ser Lys Asp Ser Pro Val Ile Gln 1235 1240 1245
- Val Leu Ala Tyr Asp Ala Asp Glu Gly Gln Asn Ala Asp Val Thr Tyr 1250 1260
- Ser Val Asn Pro Glu Asp Leu Val Lys Asp Val Ile Glu Ile Asn Pro 1265 1270 1275 1280
- Val Thr Gly Val Val Lys Val Lys Asp Ser Leu Val Gly Leu Glu Asn 1285 1290 1295
- Gln Thr Leu Asp Phe Phe Ile Lys Ala Gln Asp Gly Gly Pro Pro His 1300 1305 1310
- Trp Asn Ser Leu Val Pro Val Arg Leu Gln Val Val Pro Lys Lys Val
  1315 1320 1325
- Ser Leu Pro Lys Phe Ser Glu Pro Leu Tyr Thr Phe Ser Ala Pro Glu 1330 1335 1340
- Asp Leu Pro Glu Gly Ser Glu Ile Gly Ile Val Lys Ala Val Ala Ala 1345 1350 1355 1360

- Gln Asp Pro Val Ile Tyr Ser Leu Val Arg Gly Thr Thr Pro Glu Ser 1365 1370 1375
- Asn Lys Asp Gly Val Phe Ser Leu Asp Pro Asp Thr Gly Val Ile Lys 1380 1385 1390
- Val Arg Lys Pro Met Asp His Glu Ser Thr Lys Leu Tyr Gln Ile Asp 1395 1400 1405
- Val Met Ala His Cys Leu Gln Asn Thr Asp Val Val Ser Leu Val Ser 1410 1415 1420
- Val Asn Ile Gln Val Gly Asp Val Asn Asp Asn Arg Pro Val Phe Glu 1425 1430 1435 1440
- Ala Asp Pro Tyr Lys Ala Val Leu Thr Glu Asn Met Pro Val Gly Thr
  1445 1450 1455
- Ser Val Ile Gln Val Thr Ala Ile Asp Lys Asp Thr Gly Arg Asp Gly
  1460 1465 1470
- Gln Val Ser Tyr Arg Leu Ser Ala Asp Pro Gly Ser Asn Val His Glu 1475 1480 1485
- Leu Phe Ala Ile Asp Ser Glu Ser Gly Trp Ile Thr Thr Leu Gln Glu 1490 1495 1500
- Leu Asp Cys Glu Thr Cys Gln Thr Tyr His Phe His Val Val Ala Tyr 1505 1510 1515 1520
- Asp His Gly Gln Thr Ile Gln Leu Ser Ser Gln Ala Leu Val Gln Val 1525 1530 1535
- Ser Ile Thr Asp Glu Asn Asp Asn Ala Pro Arg Phe Ala Ser Glu Glu 1540 1545 1550
- Tyr Arg Gly Ser Val Val Glu Asn Ser Glu Pro Gly Glu Leu Val Ala 1555 1560 1565
- Thr Leu Lys Thr Leu Asp Ala Asp Ile Ser Glu Gln Asn Arg Gln Val 1570 1575 1580
- Thr Cys Tyr Ile Thr Glu Gly Asp Pro Leu Gly Gln Phe Gly Ile Ser 1585 1590 1595 1600
- Gln Val Gly Asp Glu Trp Arg Ile Ser Ser Arg Lys Thr Leu Asp Arg 1605 1610 1615
- Glu His Thr Ala Lys Tyr Leu Leu Arg Val Thr Ala Ser Asp Gly Lys
  1620 1625 1630
- Phe Gln Ala Ser Val Thr Val Glu Ile Phe Val Leu Asp Val Asn Asp 1635 1640 1645
- Asn Ser Pro Gln Cys Ser Gln Leu Leu Tyr Thr Gly Lys Val His Glu 1650 1655 1660

- Asp Val Phe Pro Gly His Phe Ile Leu Lys Val Ser Ala Thr Asp Leu 1665 1670 1680
- Asp Thr Asp Thr Asn Ala Gln Ile Thr Tyr Ser Leu His Gly Pro Gly
  1685 1690 1695
- Ala His Glu Phe Lys Leu Asp Pro His Thr Gly Glu Leu Thr Thr Leu 1700 1705 1710
- Thr Ala Leu Asp Arg Glu Arg Lys Asp Val Phe Asn Leu Val Ala Lys 1715 1720 1725
- Ala Thr Asp Gly Gly Arg Ser Cys Gln Ala Asp Ile Thr Leu His 1730 1735 1740
- Val Glu Asp Val Asn Asp Asn Ala Pro Arg Phe Phe Pro Ser His Cys 1745 1750 1755 . 1760
- Ala Val Ala Val Phe Asp Asn Thr Thr Val Lys Thr Pro Val Ala Val 1765 1770 1775
- Val Phe Ala Arg Asp Pro Asp Gln Gly Ala Asn Ala Gln Val Tyr 1780 1785 1790
- Ser Leu Pro Asp Ser Ala Glu Gly His Phe Ser Ile Asp Ala Thr Thr 1795 1800 1805
- Gly Val Ile Arg Leu Glu Lys Pro Leu Gln Val Arg Pro Gln Ala Pro 1810 1815 1820
- Leu Glu Leu Thr Val Arg Ala Ser Asp Leu Gly Thr Pro Ile Pro Leu 1825 1830 1835
- Ser Thr Leu Gly Thr Val Thr Val Ser Val Val Gly Leu Glu Asp Tyr 1845 1850 1855
- Leu Pro Val Phe Leu Asn Thr Glu His Ser Val Gln Val Pro Glu Asp 1860 1865 1870
- Ala Pro Pro Gly Thr Glu Val Leu Gln Leu Ala Thr Leu Thr Arg Pro 1875 1880 1885
- Gly Ala Glu Lys Thr Gly Tyr Arg Val Val Ser Gly Asn Glu Gln Gly 1890 1895 1900
- Arg Phe Arg Leu Asp Ala Arg Thr Gly Ile Leu Tyr Val Asn Ala Ser 1905 1910 1915 1920
- Leu Asp Phe Glu Thr Ser Pro Lys Tyr Phe Leu Ser Ile Glu Cys Ser 1925 1930 1935
- Arg Lys Ser Ser Ser Ser Leu Ser Asp Val Thr Thr Val Met Val Asn 1940 1945 1950
- Ile Thr Asp Val Asn Glu His Arg Pro Gln Phe Pro Gln Asp Pro Tyr
  1955 1960 1965

- Ser Thr Arg Val Leu Glu Asn Ala Leu Val Gly Asp Val Ile Leu Thr 1970 1975 1980
- Val Ser Ala Thr Asp Glu Asp Gly Pro Leu Asn Ser Asp Ile Thr Tyr 1985 1990 1995 2000
- Ser Leu Ile Gly Gly Asn Gln Leu Gly His Phe Thr Ile His Pro Lys  $2005 \hspace{1cm} 2010 \hspace{1cm} 2015$
- Lys Gly Glu Leu Gln Val Ala Lys Ala Leu Asp Arg Glu Gln Ala Ser 2020 2025 2030
- Ser Tyr Ser Leu Lys Leu Arg Ala Thr Asp Ser Gly Gln Pro Pro Leu 2035 2040 2045
- His Glu Asp Thr Asp Ile Ala Ile Gln Val Ala Asp Val Asn Asp Asn 2050 2055 2060
- Pro Pro Arg Phe Phe Gln Leu Asn Tyr Ser Thr Thr Val Gln Glu Asn 2065 2070 2075 2080
- Ser Pro Ile Gly Ser Lys Val Leu Gln Leu Ile Leu Ser Asp Pro Asp 2085 2090 2095
- Ser Pro Glu Asn Gly Pro Pro Tyr Ser Phe Arg Ile Thr Lys Gly Asn 2100 2105 2110
- Asn Gly Ser Ala Phe Arg Val Thr Pro Asp Gly Trp Leu Val Thr Ala 2115 2120 2125
- Glu Gly Leu Ser Arg Arg Ala Gln Glu Trp Tyr Gln Leu Gln Ile Gln 2130 2135 2140
- Ala Ser Asp Ser Gly Ile Pro Pro Leu Ser Ser Leu Thr Ser Val Arg 2145 2150 2155 2160
- Val His Val Thr Glu Gln Ser His Tyr Ala Pro Ser Ala Leu Pro Leu 2165 2170 2175
- Glu Ile Phe Ile Thr Val Gly Glu Asp Glu Phe Gln Gly Gly Met Val 2180 2185 2190
- Gly Lys Ile His Ala Thr Asp Arg Asp Pro Gln Asp Thr Leu Thr Tyr 2195 2200 2205
- Ser Leu Ala Glu Glu Glu Thr Leu Gly Arg His Phe Ser Val Gly Ala 2210 2215 2220
- Pro Asp Gly Lys Ile Ile Ala Ala Gln Gly Leu Pro Arg Gly His Tyr 2225 2230 2235 2240
- Ser Phe Asn Val Thr Val Ser Asp Gly Thr Phe Thr Thr Thr Ala Gly 2245 2250 2255
- Val His Val Tyr Val Trp His Val Gly Gln Glu Ala Leu Gln Gln Ala 2260 2265 2270

- Met Trp Met Gly Phe Tyr Gln Leu Thr Pro Glu Glu Leu Val Ser Asp 2275 2280 2285
- His Trp Arg Asn Leu Gln Arg Phe Leu Ser His Lys Leu Asp Ile Lys 2290 2295 2300
- Arg Ala Asn Ile His Leu Ala Ser Leu Gln Pro Ala Glu Ala Val Ala 2305 2310 2315 2320
- Gly Val Asp Val Leu Leu Val Phe Glu Gly His Ser Gly Thr Phe Tyr 2325 2330 2335
- Glu Phe Gln Glu Leu Ala Ser Ile Ile Thr His Ser Ala Lys Glu Met 2340 2345 2350
- Glu His Ser Val Gly Val Gln Met Arg Ser Ala Met Pro Met Val Pro 2355 2360 2365
- Cys Gln Gly Pro Thr Cys Gln Gly Gln Ile Cys His Asn Thr Val His 2370 2375 2380
- Leu Asp Pro Lys Val Gly Pro Thr Tyr Ser Thr Ala Arg Leu Ser Ile 2385 2390 2395 2400
- Leu Thr Pro Arg His His Leu Gln Arg Ser Cys Ser Cys Asn Gly Thr 2405 2410 2415
- Ala Thr Arg Phe Ser Gly Gln Ser Tyr Val Arg Tyr Arg Ala Pro Ala 2420 2425 2430
- Ala Arg Asn Trp His Ile His Phe Tyr Leu Lys Thr Leu Gln Pro Gln 2435 2440 2445
- Ala Ile Leu Leu Phe Thr Asn Glu Thr Ala Ser Val Ser Leu Lys Leu 2450 2455 2460
- Ala Ser Gly Val Pro Gln Leu Glu Tyr His Cys Leu Gly Gly Phe Tyr 2465 2470 2475 2480
- Gly Asn Leu Ser Ser Gln Arg His Val Asn Asp His Glu Trp His Ser 2485 2490 2495
- Ile Leu Val Glu Glu Met Asp Ala Ser Ile Arg Leu Met Val Asp Ser 2500 2505 2510
- Met Gly Asn Thr Ser Leu Val Val Pro Glu Asn Cys Arg Gly Leu Arg 2515 2520 2525
- Pro Glu Arg His Leu Leu Leu Gly Gly Leu Ile Leu Leu His Ser Ser 2530 2540
- Ser Asn Val Ser Gln Gly Phe Glu Gly Cys Leu Asp Ala Val Val 2545 2550 2555 2560
- Asn Glu Glu Ala Leu Asp Leu Leu Ala Pro Gly Lys Thr Val Ala Gly
  2565 2570 2575

- Leu Leu Glu Thr Gln Ala Leu Thr Gln Cys Cys Leu His Ser Asp Tyr 2580 2585 2590
- Cys Ser Gln Asn Thr Cys Leu Asn Gly Gly Lys Cys Ser Trp Thr His 2595 2600 2605
- Gly Ala Gly Tyr Val Cys Lys Cys Pro Pro Gln Phe Ser Gly Lys His 2610 2615 2620
- Cys Glu Gln Gly Arg Glu Asn Cys Thr Phe Ala Pro Cys Leu Glu Gly 2625 2630 2635 2640
- Gly Thr Cys Ile Leu Ser Pro Lys Gly Ala Ser Cys Asn Cys Pro His 2645 2650 2655
- Pro Tyr Thr Gly Asp Arg Cys Glu Met Glu Ala Arg Gly Cys Ser Glu 2660 2665 2670
- Gly His Cys Leu Val Thr Pro Glu Ile Gln Arg Gly Asp Trp Gly Gln 2675 2680 2685
- Gln Glu Leu Leu Ile Ile Thr Val Ala Val Ala Phe Ile Ile Ser 2690 2695 2700
- Thr Val Gly Leu Leu Phe Tyr Cys Arg Arg Cys Lys Ser His Lys Pro 2705 2710 2715 2720
- Val Ala Met Glu Asp Pro Asp Leu Leu Ala Arg Ser Val Gly Val Asp 2725 2730 2735
- Thr Gln Ala Met Pro Ala Ile Glu Leu Asn Pro Leu Ser Ala Ser Ser 2740 2745 2750
- Cys Asn Asn Leu Asn Gln Pro Glu Pro Ser Lys Ala Ser Val Pro Asn 2755 2760 2765
- Glu Leu Val Thr Phe Gly Pro Asn Ser Lys Gln Arg Pro Val Val Cys 2770 2775 2780
- Ser Val Pro Pro Arg Leu Pro Pro Ala Ala Val Pro Ser His Ser Asp 2785 2790 2795 2800
- Asn Glu Pro Val Ile Lys Arg Thr Trp Ser Ser Glu Glu Met Val Tyr 2805 2810 2815
- Pro Gly Gly Ala Met Val Trp Pro Pro Thr Tyr Ser Arg Asn Glu Arg 2820 2825 2830
- Trp Glu Tyr Pro His Ser Glu Val Thr Gln Gly Pro Leu Pro Pro Ser 2835 2840 2845
- Ala His Arg His Ser Thr Pro Val Val Met Pro Glu Pro Asn Gly Leu 2850 2855 2860
- Tyr Gly Gly Phe Pro Phe Pro Leu Glu Met Glu Asn Lys Arg Ala Pro 2865 2870 2875 2880

- Leu Pro Pro Arg Tyr Ser Asn Gln Asn Leu Glu Asp Leu Met Pro Ser 2885 2890 2895
- Arg Pro Pro Ser Pro Arg Glu Arg Leu Val Ala Pro Cys Leu Asn Glu 2900 2905 2910
- Tyr Thr Ala Ile Ser Tyr Tyr His Ser Gln Phe Arg Gln Gly Gly Gly 2915 2920 2925
- Gly Pro Cys Leu Ala Asp Gly Gly Tyr Lys Gly Val Gly Met Arg Leu 2930 2935 2940
- Ser Arg Ala Gly Pro Ser Tyr Ala Val Cys Glu Val Glu Gly Ala Pro 2945 2950 2955 2960
- Leu Ala Gly Gln Gly Gln Pro Arg Val Pro Pro Asn Tyr Glu Gly Ser 2965 2970 2975
- Asp Met Val Glu Ser Asp Tyr Gly Ser Cys Glu Glu Val Met Phe 2980 2985 2990
- <210> 23
- <211> 928
- <212> PRT
- <213> Homo sapiens
- <400> 23
- His Cys Gln Glu Asp Gly Ile Met Leu Ser Ala Asp Cys Ser Glu Leu 1 5 10 15
- Gly Leu Ser Ala Val Pro Gly Asp Leu Asp Pro Leu Thr Ala Tyr Leu
  20 25 30
- Asp Leu Ser Met Asn Asn Leu Thr Glu Leu Gln Pro Gly Leu Phe His
  35 40 45
- His Leu Arg Phe Leu Glu Glu Leu Arg Leu Ser Gly Asn His Leu Ser 50 55 60
- His Ile Pro Gly Gln Ala Phe Ser Gly Leu Tyr Ser Leu Lys Ile Leu 65 70 75 80
- Met Leu Gln Asn Asn Gln Leu Gly Gly Ile Pro Ala Glu Ala Leu Trp 85 90 95
- Glu Leu Pro Ser Leu Gln Ser Leu Arg Leu Asp Ala Asn Leu Ile Ser 100 105 110
- Leu Val Pro Glu Arg Ser Phe Glu Gly Leu Ser Ser Leu Arg His Leu 115 120 125
- Trp Leu Asp Asp Asn Ala Leu Thr Glu Ile Pro Val Arg Ala Leu Asn 130 135 140
- Asn Leu Pro Ala Leu Gln Ala Met Thr Leu Ala Leu Asn Arg Ile Ser 145 150 155 160

His Ile Pro Asp Tyr Ala Phe Gln Asn Leu Thr Ser Leu Val Val Leu 170 His Leu His Asn Asn Arg Ile Gln His Leu Gly Thr His Ser Phe Glu 185 Gly Leu His Asn Leu Glu Thr Leu Asp Leu Asn Tyr Asn Lys Leu Gln Glu Phe Pro Val Ala Ile Arg Thr Leu Gly Arg Leu Gln Glu Leu Gly 215 Phe His Asn Asn Asn Ile Lys Ala Ile Pro Glu Lys Ala Phe Met Gly 230 235 Asn Pro Leu Leu Gln Thr Ile His Phe Tyr Asp Asn Pro Ile Gln Phe 245 250 Val Gly Arg Ser Ala Phe Gln Tyr Leu Pro Lys Leu His Thr Leu Ser 265 Leu Asn Gly Ala Met Asp Ile Gln Glu Phe Pro Asp Leu Lys Gly Thr Thr Ser Leu Glu Ile Leu Thr Leu Thr Arg Ala Gly Ile Arg Leu Leu 295 Pro Ser Gly Met Cys Gln Gln Leu Pro Arg Leu Arg Val Leu Glu Leu 310 315 Ser His Asn Gln Ile Glu Glu Leu Pro Ser Leu His Arq Cys Gln Lys 330 Leu Glu Glu Ile Gly Leu Gln His Asn Arg Ile Trp Glu Ile Gly Ala Asp Thr Phe Ser Gln Leu Ser Ser Leu Gln Ala Leu Asp Leu Ser Trp 355 Asn Ala Ile Arg Ser Ile His Pro Glu Ala Phe Ser Thr Leu His Ser 375 Leu Val Lys Leu Asp Leu Thr Asp Asn Gln Leu Thr Thr Leu Pro Leu 390 395 Ala Gly Leu Gly Gly Leu Met His Leu Lys Leu Lys Gly Asn Leu Ala Leu Ser Gln Ala Phe Ser Lys Asp Ser Phe Pro Lys Leu Arg Ile Leu Glu Val Pro Tyr Ala Tyr Gln Cys Cys Pro Tyr Gly Met Cys Ala Ser 435 Phe Phe Lys Ala Ser Gly Gln Trp Glu Ala Glu Asp Leu His Leu Asp 450 455 460

Asp Glu Glu Ser Ser Lys Arg Pro Leu Gly Leu Leu Ala Arg Gln Ala 470 475 Glu Asn His Tyr Asp Gln Asp Leu Asp Glu Leu Gln Leu Glu Met Glu 490 Asp Ser Lys Pro His Pro Ser Val Gln Cys Ser Pro Thr Pro Gly Pro 500 Phe Lys Pro Cys Glu Tyr Leu Phe Glu Ser Trp Gly Ile Arg Leu Ala 520 Val Trp Ala Ile Val Leu Leu Ser Val Leu Cys Asn Gly Leu Val Leu 535 Leu Thr Val Phe Ala Gly Gly Pro Ala Pro Leu Pro Pro Val Lys Phe 550 555 Val Val Gly Ala Ile Ala Gly Ala Asn Thr Leu Thr Gly Ile Ser Cys 570 Gly Leu Leu Ala Ser Val Asp Ala Leu Thr Phe Gly Gln Phe Ser Glu 580 Tyr Gly Ala Arg Trp Glu Thr Gly Leu Gly Cys Arg Ala Thr Gly Phe 600 Leu Ala Val Leu Gly Ser Glu Ala Ser Val Leu Leu Leu Thr Leu Ala Ala Val Gln Cys Ser Val Ser Val Ser Cys Val Arg Ala Tyr Gly Lys Ser Pro Ser Leu Gly Ser Val Arg Ala Gly Val Leu Gly Cys Leu Ala 650 Leu Ala Gly Leu Ala Ala Leu Pro Leu Ala Ser Val Gly Glu Tyr 660 Gly Ala Ser Pro Leu Cys Leu Pro Tyr Ala Pro Pro Glu Gly Gln Pro 680 Ala Ala Leu Gly Phe Thr Val Ala Leu Val Met Met Asn Ser Phe Cys 690 Phe Leu Val Val Ala Gly Ala Tyr Ile Lys Leu Tyr Cys Asp Leu Pro 705 Arg Gly Asp Phe Glu Ala Val Trp Asp Cys Ala Met Val Arg His Val 725 730 Ala Trp Leu Ile Phe Ala Asp Gly Leu Leu Tyr Cys Pro Val Ala Phe 740 Leu Ser Phe Ala Ser Met Leu Gly Leu Phe Pro Val Thr Pro Glu Ala 755 760 765

Val Lys Ser Val Leu Leu Val Val Leu Pro Leu Pro Ala Cys Leu Asn 775 Pro Leu Leu Tyr Leu Leu Phe Asn Pro His Phe Arg Asp Leu Arg 790 795 Arg Leu Arg Pro Arg Ala Gly Asp Ser Gly Pro Leu Ala Tyr Ala Ala Ala Gly Glu Leu Glu Lys Ser Ser Cys Asp Ser Thr Gln Ala Leu Val 825 Ala Phe Ser Asp Val Asp Leu Ile Leu Glu Ala Ser Glu Ala Gly Arq 840 Pro Pro Gly Leu Glu Thr Tyr Gly Phe Pro Ser Val Thr Leu Ile Ser 855 Cys Gln Gln Pro Gly Ala Pro Arg Leu Glu Gly Ser His Cys Val Glu 870 Pro Glu Gly Asn His Phe Gly Asn Pro Gln Pro Ser Met Asp Gly Glu Leu Leu Arg Ala Glu Gly Ser Thr Pro Ala Gly Gly Leu Ser

905

Gly Gly Gly Phe Gln Pro Ser Gly Leu Ala Phe Ala Ser His Val 915 920 925

<210> 24

<211> 893

<212> PRT

<213> Homo sapiens

<400> 24

Met Asn Asn Leu Thr Glu Leu Gln Pro Gly Leu Phe His His Leu Arg

1 10 15

Phe Leu Glu Glu Leu Arg Leu Ser Gly Asn His Leu Ser His Ile Pro 20 25 30

Gly Gln Ala Phe Ser Gly Leu Tyr Ser Leu Lys Ile Leu Met Leu Gln 35 40 45

Asn Asn Gln Leu Gly Gly Ile Pro Ala Glu Ala Leu Trp Glu Leu Pro 50 55 60

Ser Leu Gln Ser Leu Arg Leu Asp Ala Asn Leu Ile Ser Leu Val Pro 65 70 75 80

Glu Arg Ser Phe Glu Gly Leu Ser Ser Leu Arg His Leu Trp Leu Asp

Asp	Asn	Ala	Leu 100	Thr	Glu	Ile	Pro	Val 105	Arg	Ala	Leu	Asn	Asn 110	Leu	Pro
Ala	Leu	Gln 115		Met	Thr	Leu	Ala 120	Leu	Asn	Arg	Ile	Ser 125	His	Ile	Pro
Asp	Tyr 130	Ala	Phe	Gln	Asn	Leu 135	Thr	Ser	Leu	Val	Val 140	Leu	His	Leu	His
Asn 145		Arg	Ile	Gln	His 150	Leu	Gly	Thr	His	Ser 155	Phe	Glu	Gly	Leu	His 160
Asn	Leu	Glu	Thr	Leu 165	Asp	Leu	Asn	Tyr	Asn 170	Lys	Leu	Gln	Glu	Phe 175	Pro
Val	Ala	Ile	Arg 180	Thr	Leu	Gly	Arg	Leu 185	Gln	Glu	Leu	Gly	Phe 190	His	Asr
Asn	Asn	Ile 195	Lys	Ala	Ile	Pro	Glu 200	Lys	Ala	Phe	Met	Gly 205	Asn	Pro	Leı
Leu	Gln 210	Thr	Ile	His	Phe	Tyr 215	Asp	Asn	Pro	Ile	Gln 220	Phe	Val	Gly	Arg
Ser 225	Ala	Phe	Gln	Tyr	Leu 230	Pro	Lys	Leu	His	Thr 235	Leu	Ser	Leu	Asn	Gly 240
Ala	Met	Asp	Ile	Gln 245	Glu	Phe	Pro	Asp	Leu 250	Lys	Gly	Thr	Thr	Ser 255	Leu
Glu	Ile	Leu	Thr 260	Leu	Thr	Arg	Ala	Gly 265	Ile	Arg	Leu	Leu	Pro 270	Ser	Gly
Met	Cys	Gln 275	Gln	Leu	Pro	Arg	Leu 280	Arg	Val	Leu	Glu	Leu 285	Ser	His	Asn
Gln	Ile 290	Glu	Glu	Leu	Pro	Ser 295	Leu	His	Arg	Cys	Gln 300	Lys	Leu	Glu	Glu
Ile 305	Gly	Leu	Gln	His	Asn 310	Arg	Ile	Trp	Glu	Ile 315	Gly	Ala	Asp	Thr	Phe 320
Ser	Gln	Leu	Ser	Ser 325	Leu	Gln	Ala	Leu	Asp 330	Leu	Ser	Trp	Asn	Ala 335	Ile
Arg	Ser	Ile	His 340	Pro	Glu	Ala	Phe	Ser 345	Thr	Leu	His	Ser	Leu 350	Val	Lys
Leu	Asp	Leu 355	Thr	Asp	Asn	Gln	Leu 360	Thr	Thr	Leu	Pro	Leu 365	Ala	Gly	Leu
Gly	Gly 370	Leu	Met	His	Leu	Lys 375	Leu	Lys	Gly	Asn	Leu 380	Ala	Leu	Ser	Gln

Ala Phe Ser Lys Asp Ser Phe Pro Lys Leu Arg Ile Leu Glu Val Pro

385 390	395 400
---------	---------

- Tyr Ala Tyr Gln Cys Cys Pro Tyr Gly Met Cys Ala Ser Phe Phe Lys 405 410 415
- Ala Ser Gly Gln Trp Glu Ala Glu Asp Leu His Leu Asp Asp Glu Glu
  420 425 430
- Ser Ser Lys Arg Pro Leu Gly Leu Leu Ala Arg Gln Ala Glu Asn His
  435
  440
  445
- Tyr Asp Gln Asp Leu Asp Glu Leu Gln Leu Glu Met Glu Asp Ser Lys 450 455 460
- Pro His Pro Ser Val Gln Cys Ser Pro Thr Pro Gly Pro Phe Lys Pro 465 470 475 480
- Cys Glu Tyr Leu Phe Glu Ser Trp Gly Ile Arg Leu Ala Val Trp Ala 485 490 495
- Ile Val Leu Ser Val Leu Cys Asn Gly Leu Val Leu Leu Thr Val
  500 505 510
- Phe Ala Gly Gly Pro Val Pro Leu Pro Pro Val Lys Phe Val Val Gly 515 520 525
- Ala Ile Ala Gly Ala Asn Thr Leu Thr Gly Ile Ser Cys Gly Leu Leu 530 535 540
- Ala Ser Val Asp Ala Leu Thr Phe Gly Gln Phe Ser Glu Tyr Gly Ala 545 550 560
- Arg Trp Glu Thr Gly Leu Gly Cys Arg Ala Thr Gly Phe Leu Ala Val
  565 570 575
- Leu Gly Ser Glu Ala Ser Val Leu Leu Leu Thr Leu Ala Ala Val Gln 580 585 590
- Cys Ser Val Ser Val Ser Cys Val Arg Ala Tyr Gly Lys Ser Pro Ser 595 600 605
- Leu Gly Ser Val Arg Ala Gly Val Leu Gly Cys Leu Ala Leu Ala Gly 610 615 620
- Leu Ala Ala Leu Pro Leu Ala Ser Val Gly Glu Tyr Gly Ala Ser 625 630 635 640
- Pro Leu Cys Leu Pro Tyr Ala Pro Pro Glu Gly Gln Pro Ala Ala Leu 645. 650 655
- Gly Phe Thr Val Ala Leu Val Met Met Asn Ser Phe Cys Phe Leu Val 660 665 670
- Val Ala Gly Ala Tyr Ile Lys Leu Tyr Cys Asp Leu Pro Arg Gly Asp 675 680 685
- Phe Glu Ala Val Trp Asp Cys Ala Met Val Arg His Val Ala Trp Leu

Ile Phe Ala Asp GlyLeu Leu Tyr CysPro Val Ala Phe Leu Ser Phe705710715720Ala Ser Met Leu GlyLeu Phe Pro Val Thr Pro Glu Ala Val LysSer725730735

Val Leu Leu Val Val Leu Pro Leu Pro Ala Cys Leu Asn Pro Leu Leu
740 745 750

Tyr Leu Leu Phe Asn Pro His Phe Arg Asp Asp Leu Arg Arg Leu Arg
755 760 765

Pro Arg Ala Gly Asp Ser Gly Pro Leu Ala Tyr Ala Ala Ala Gly Glu 770 780

Leu Glu Lys Ser Ser Cys Asp Ser Thr Gln Ala Leu Val Ala Phe Ser 785 790 795 800

Asp Val Asp Leu Ile Leu Glu Ala Ser Glu Ala Gly Arg Pro Pro Gly 805 810 815

Leu Glu Thr Tyr Gly Phe Pro Ser Val Thr Leu Ile Ser Cys Gln Gln 820 825 830

Pro Gly Ala Pro Arg Leu Glu Gly Ser His Cys Val Glu Pro Glu Gly 835 840 845

Asn His Phe Gly Asn Pro Gln Pro Ser Met Asp Gly Glu Leu Leu 850 855 860

Arg Ala Glu Gly Ser Thr Pro Ala Gly Gly Gly Leu Ser Gly Gly 865 870 875 880

Gly Phe Gln Pro Ser Gly Leu Ala Phe Ala Ser His Val 885 890

<210> 25

<211> 828

<212> PRT

<213> Homo sapiens

<400> 25

Met Arg Leu Glu Gly Glu Gly Arg Ser Ala Arg Ala Gly Gln Asn Leu

1 10 15

Ser Arg Ala Gly Ser Ala Arg Arg Gly Ala Pro Arg Asp Leu Ser Met
20 25 30

Asn Asn Leu Thr Glu Leu Gln Pro Gly Leu Phe His His Leu Arg Phe
35 40 45

Leu Glu Glu Leu Arg Leu Ser Gly Asn His Leu Ser His Ile Pro Gly
50 55 60

Gln 65		Phe	Ser	Gly	Leu 70	Tyr	Ser	Leu	Lys	Ile 75		Met	Leu	Gln	Asn 80
Asn	Gln	Leu	Gly	Gly 85	Ile	Pro	Ala	Glu	Ala 90	Leu	Trp	Glu	Leu	Pro 95	Ser
Leu	Gln	Ser	Leu 100	Asp	Leu	Asn	Tyr	Asn 105	_	Leu	Gln	Glu	Phe 110	Pro	Val
Ala	Ile	Arg 115	Thr	Leu	Gly	Arg	Leu 120	Gln	Glu	Leu	Gly	Phe 125	His	Asn	Asn
Asn	Ile 130	Lys	Ala	Ile	Pro	Glu 135	Lys	Ala	Phe	Met	Gly 140	Asn	Pro	Leu	Leu
Gln 145	Thr	Ile	His	Phe	Tyr 150	Asp	Asn	Pro	Ile	Gln 155	Phe	Val	Gly	Arg	Ser 160
Ala	Phe	Gln	Tyr	Leu 165	Pro	Lys	Leu	His	Thr 170	Leu	Ser	Leu	Asn	Gly 175	Ala
Met	Asp	Ile	Gln 180	Glu	Phe	Pro	Asp	Leu 185	Lys	Gly	Thr	Thr	Ser 190	Leu	Glu
Ile	Leu	Thr 195	Leu	Thr	Arg	Ala	Gly 200	Ile	Arg	Leu	Leu	Pro 205	Ser	Gly	Met
Cys	Gln 210	Gln	Leu	Pro	Arg	Leu 215	Arg	Val	Leu	Glu	Leu 220	Ser	His	Asn	Gln
Ile 225	Glu	Glu	Leu	Pro	Ser 230	Leu	His	Arg	Суѕ	Gln 235	Lys	Leu	Glu	Glu	Ile 240
Gly	Leu	Gln	His	Asn 245	Arg	Ile	Trp	Glu	Ile 250	Gly	Ala	Asp	Thr	Phe 255	Ser
Gln	Leu	Ser	Ser 260	Leu	Gln	Ala	Leu	Asp 265	Leu	Ser	Trp	Asn	Ala 270	Ile	Arg
Ser	Ile	His 275	Pro	Glu	Ala	Phe	Ser 280	Thr	Leu	His	Ser	Leu 285	Val	Lys	Leu
Asp	Leu 290	Thr	Asp	Asn	Gln	Leu 295	Thr	Thr	Leu	Pro	Leu 300	Ala	Gly	Leu	Gly
Gly 305	Leu	Met	His	Leu	Lys 310	Leu	Lys	Gly	Asn	Leu 315	Ala	Leu	Ser	Gln	Ala 320
Phe	Ser	Lys	Asp	Ser 325	Phe	Pro	Lys	Leu	Arg 330	Ile	Leu	Glu	Val	Pro 335	Tyr
Ala	Tyr	Gln	Cys 340	Cys	Pro	Tyr	Gly	Met 345	Cys	Ala	Ser	Phe	Phe 350	Lys	Ala
Ser	Gly	Gln	Trp	Glu	Ala		Asp		His	Leu	Asp	Asp	Glu	Glu	Ser

Ser Lys Arg Pro Leu Gly Leu Leu Ala Arg Gln Ala Glu Asn His Tyr 370 Asp Gln Asp Leu Asp Glu Leu Gln Leu Glu Met Glu Asp Ser Lys Pro 390 His Pro Ser Val Gln Cys Ser Pro Thr Pro Gly Pro Phe Lys Pro Cys 405 410 Glu Tyr Leu Phe Glu Ser Trp Gly Ile Arg Leu Ala Val Trp Ala Ile 420 425 Val Leu Leu Ser Val Leu Cys Asn Gly Leu Val Leu Leu Thr Val Phe Ala Gly Gly Pro Val Pro Leu Pro Pro Val Lys Phe Val Val Gly Ala 455 Ile Ala Gly Ala Asn Thr Leu Thr Gly Ile Ser Cys Gly Leu Leu Ala 470 Ser Val Asp Ala Leu Thr Phe Gly Gln Phe Ser Glu Tyr Gly Ala Arg 490 Trp Glu Thr Gly Leu Gly Cys Arg Ala Thr Gly Phe Leu Ala Val Leu 505 Gly Ser Glu Ala Ser Val Leu Leu Leu Thr Leu Ala Ala Val Gln Cys Ser Val Ser Val Ser Cys Val Arg Ala Tyr Gly Lys Ser Pro Ser Leu Gly Ser Val Arg Ala Gly Val Leu Gly Cys Leu Ala Leu Ala Gly Leu 545 550 Ala Ala Leu Pro Leu Ala Ser Val Gly Glu Tyr Gly Ala Ser Pro 570 Leu Cys Leu Pro Tyr Ala Pro Pro Glu Gly Gln Pro Ala Ala Leu Gly 580 Phe Thr Val Ala Leu Val Met Met Asn Ser Phe Cys Phe Leu Val Val 600 Ala Gly Ala Tyr Ile Lys Leu Tyr Cys Asp Leu Pro Arg Gly Asp Phe 615 Glu Ala Val Trp Asp Cys Ala Met Val Arg His Val Ala Trp Leu Ile 625 630 635 640 Phe Ala Asp Gly Leu Leu Tyr Cys Pro Val Ala Phe Leu Ser Phe Ala 645 650 Ser Met Leu Gly Leu Phe Pro Val Thr Pro Glu Ala Val Lys Ser Val 665 670

- Leu Leu Val Val Leu Pro Leu Pro Ala Cys Leu Asn Pro Leu Leu Tyr 675 680 685
- Leu Leu Phe Asn Pro His Phe Arg Asp Asp Leu Arg Arg Leu Arg Pro 690 695 700
- Arg Ala Gly Asp Ser Gly Pro Leu Ala Tyr Ala Ala Ala Gly Glu Leu 705 710 715 720
- Glu Lys Ser Ser Cys Asp Ser Thr Gln Ala Leu Val Ala Phe Ser Asp
  725 730 735
- Val Asp Leu Ile Leu Glu Ala Ser Glu Ala Gly Arg Pro Pro Gly Leu
  740 745 750
- Glu Thr Tyr Gly Phe Pro Ser Val Thr Leu Ile Ser Cys Gln Gln Pro 755 760 765
- Gly Ala Pro Arg Leu Glu Gly Ser His Cys Val Glu Pro Glu Gly Asn 770 775 780
- His Phe Gly Asn Pro Gln Pro Ser Met Asp Gly Glu Leu Leu Leu Arg
  785 790 795 800
- Ala Glu Gly Ser Thr Pro Ala Gly Gly Gly Leu Ser Gly Gly Gly 805 810 815
- Phe Gln Pro Ser Gly Leu Ala Leu Leu His Thr Tyr 820 825

<210> 26

<211> 907

<212> PRT

<213> Homo sapiens

<400> 26

- Met Asp Thr Ser Arg Leu Gly Val Leu Leu Ser Leu Pro Val Leu Leu 1 5 10 15
- Gln Leu Ala Thr Gly Gly Ser Ser Pro Arg Ser Gly Val Leu Leu Arg
  20 25 30
- Gly Cys Pro Thr His Cys His Cys Glu Pro Asp Gly Arg Met Leu Leu 35 40 45
- Arg Val Asp Cys Ser Asp Leu Gly Leu Ser Glu Leu Pro Ser Asn Leu 50 55 60
- Ser Val Phe Thr Ser Tyr Leu Asp Leu Ser Met Asn Asn Ile Ser Gln 65 70 75 80
- Leu Leu Pro Asn Pro Leu Pro Ser Leu Arg Phe Leu Glu Glu Leu Arg
  85 90 95
- Leu Ala Gly Asn Ala Leu Thr Tyr Ile Pro Lys Gly Ala Phe Thr Gly
  100 105 110

Leu Tyr Ser Leu Lys Val Leu Met Leu Gln Asn Asn Gln Leu Arg His 120 Val Pro Thr Glu Ala Leu Gln Asn Leu Arg Ser Leu Gln Ser Leu Arg Leu Asp Ala Asn His Ile Ser Tyr Val Pro Pro Ser Cys Phe Ser Gly 150 Leu His Ser Leu Arg His Leu Trp Leu Asp Asp Asn Ala Leu Thr Glu 170 Ile Pro Val Gln Ala Phe Arg Ser Leu Ser Ala Leu Gln Ala Met Thr 185 Leu Ala Leu Asn Lys Ile His His Ile Pro Asp Tyr Ala Phe Gly Asn 200 Leu Ser Ser Leu Val Val Leu His Leu His Asn Asn Arg Ile His Ser 215 Leu Gly Lys Lys Cys Phe Asp Gly Leu His Ser Leu Glu Thr Leu Asp 225 230 Leu Asn Tyr Asn Asn Leu Asp Glu Phe Pro Thr Ala Ile Arg Thr Leu 250 Ser Asn Leu Lys Glu Leu Gly Phe His Ser Asn Asn Ile Arg Ser Ile 265 Pro Glu Lys Ala Phe Val Gly Asn Pro Ser Leu Ile Thr Ile His Phe Tyr Asp Asn Pro Ile Gln Phe Val Gly Arg Ser Ala Phe Gln His Leu 295 Pro Glu Leu Arg Thr Leu Thr Leu Asn Gly Ala Ser Gln Ile Thr Glu 305 310 Phe Pro Asp Leu Thr Gly Thr Ala Asn Leu Glu Ser Leu Thr Leu Thr 330 Gly Ala Gln Ile Ser Ser Leu Pro Gln Thr Val Cys Asn Gln Leu Pro 340 Asn Leu Gln Val Leu Asp Leu Ser Tyr Asn Leu Leu Glu Asp Leu Pro 355 Ser Phe Ser Val Cys Gln Lys Leu Gln Lys Ile Asp Leu Arg His Asn 375 Glu Ile Tyr Glu Ile Lys Val Asp Thr Phe Gln Gln Leu Leu Ser Leu 385 390 395 Arg Ser Leu Asn Leu Ala Trp Asn Lys Ile Ala Ile Ile His Pro Asn 405 410 415

Ala Phe Ser Thr Leu Pro Ser Leu Ile Lys Leu Asp Leu Ser Ser Asn 420 425 Leu Leu Ser Ser Phe Pro Ile Thr Gly Leu His Gly Leu Thr His Leu 440 Lys Leu Thr Gly Asn His Ala Leu Gln Ser Leu Ile Ser Ser Glu Asn Phe Pro Glu Leu Lys Val Ile Glu Met Pro Tyr Ala Tyr Gln Cys Cys 470 475 Ala Phe Gly Val Cys Glu Asn Ala Tyr Lys Ile Ser Asn Gln Trp Asn 490 Lys Gly Asp Asn Ser Ser Met Asp Asp Leu His Lys Lys Asp Ala Gly 500 505 Met Phe Gln Ala Gln Asp Glu Arg Asp Leu Glu Asp Phe Leu Leu Asp 520 Phe Glu Glu Asp Leu Lys Ala Leu His Ser Val Gln Cys Ser Pro Ser 530 Pro Gly Pro Phe Lys Pro Cys Glu His Leu Leu Asp Gly Trp Leu Ile Arg Ile Gly Val Trp Thr Ile Ala Val Leu Ala Leu Thr Cys Asn Ala 570 Leu Val Thr Ser Thr Val Phe Arg Ser Pro Leu Tyr Ile Ser Pro Ile Lys Leu Leu Ile Gly Val Ile Ala Ala Val Asn Met Leu Thr Gly Val 600 Ser Ser Ala Val Leu Ala Gly Val Asp Ala Phe Thr Phe Gly Ser Phe 610 Ala Arg His Gly Ala Trp Trp Glu Asn Gly Val Gly Cys His Val Ile Gly Phe Leu Ser Ile Phe Ala Ser Glu Ser Ser Val Phe Leu Leu Thr 645 650 Leu Ala Ala Leu Glu Arg Gly Phe Ser Val Lys Tyr Ser Ala Lys Phe 660 Glu Thr Lys Ala Pro Phe Ser Ser Leu Lys Val Ile Ile Leu Leu Cys 680 Ala Leu Leu Ala Leu Thr Met Ala Ala Val Pro Leu Leu Gly Gly Ser 690 Lys Tyr Gly Ala Ser Pro Leu Cys Leu Pro Leu Pro Phe Gly Glu Pro 705 710 715 720

- Ser Thr Met Gly Tyr Met Val Ala Leu Ile Leu Leu Asn Ser Leu Cys
  725 730 735
- Phe Leu Met Met Thr Ile Ala Tyr Thr Lys Leu Tyr Cys Asn Leu Asp 740 745 750
- Lys Gly Asp Leu Glu Asn Ile Trp Asp Cys Ser Met Val Lys His Ile 755 760 765
- Ala Leu Leu Phe Thr Asn Cys Ile Leu Asn Cys Pro Val Ala Phe
  770 780
- Leu Ser Phe Ser Ser Leu Ile Asn Leu Thr Phe Ile Ser Pro Glu Val 785 790 795 800
- Ile Lys Phe Ile Leu Leu Val Val Val Pro Leu Pro Ala Cys Leu Asn 805 810 815
- Pro Leu Leu Tyr Ile Leu Phe Asn Pro His Phe Lys Glu Asp Leu Val 820 825 830
- Ser Leu Arg Lys Gln Thr Tyr Val Trp Thr Arg Ser Lys His Pro Ser 835 840 845
- Leu Met Ser Ile Asn Ser Asp Asp Val Glu Lys Gln Ser Cys Asp Ser 850 855 860
- Thr Gln Ala Leu Val Thr Phe Thr Ser Ser Ser Ile Thr Tyr Asp Leu 865 870 875 880
- Pro Pro Ser Ser Val Pro Ser Pro Ala Tyr Pro Val Thr Glu Ser Cys 885 890 895
- His Leu Ser Ser Val Ala Phe Val Pro Cys Leu 900 905

<210> 27

<211> 907

<212> PRT

<213> Homo sapiens

<400> 27

- Met Asp Thr Ser Arg Leu Gly Val Leu Leu Ser Leu Pro Val Leu Leu 1 5 10 15
- Gln Leu Ala Thr Gly Gly Ser Ser Pro Arg Ser Gly Val Leu Leu Arg
  20 25 30
- Gly Cys Pro Thr His Cys His Cys Glu Pro Asp Gly Arg Met Leu Leu 35 40 45
- Arg Val Asp Cys Ser Asp Leu Gly Leu Ser Glu Leu Pro Ser Asn Leu 50 55 60
- Ser Val Phe Thr Ser Tyr Leu Asp Leu Ser Met Asn Asn Ile Ser Gln

- Leu Leu Pro Asn Pro Leu Pro Ser Leu His Phe Leu Glu Glu Leu Arg 85 90 95
- Leu Ala Gly Asn Ala Leu Thr Tyr Ile Pro Lys Gly Ala Phe Thr Gly
  100 105 110
- Leu Tyr Ser Leu Lys Val Leu Met Leu Gln Asn Asn Gln Leu Arg His 115 120 125
- Val Pro Thr Glu Ala Leu Gln Asn Leu Arg Ser Leu Gln Ser Leu Arg 130 135 140
- Leu Asp Ala Asn His Ile Ser Tyr Val Pro Pro Ser Cys Phe Ser Gly
  145 150 155 160
- Leu His Ser Leu Arg His Leu Trp Leu Asp Asp Asn Ala Leu Thr Glu 165 170 175
- Ile Pro Val Gln Ala Phe Arg Ser Leu Ser Ala Leu Gln Ala Met Thr 180 185 190
- Leu Ala Leu Asn Lys Ile His His Ile Pro Asp Tyr Ala Phe Gly Asn 195 200 205
- Leu Ser Ser Trp Val Val Leu His Leu His Asn Asn Arg Ile His Ser 210 215 220
- Leu Gly Lys Lys Cys Phe Asp Gly Leu His Ser Leu Glu Thr Leu Asp 225 230 235 240
- Leu Asn Tyr Asn Asn Leu Asp Glu Phe Pro Thr Ala Ile Arg Thr Leu 245 250 255
- Ser Asn Leu Lys Glu Leu Gly Phe His Ser Asn Asn Ile Arg Ser Ile 260 265 270
- Pro Glu Lys Ala Phe Val Gly Asn Pro Ser Leu Ile Thr Ile His Phe 275 280 285
- Tyr Asp Asn Pro Ile Gln Phe Val Gly Arg Ser Ala Phe Gln His Leu 290 295 300
- Pro Glu Leu Arg Thr Leu Thr Leu Asn Gly Ala Ser Gln Ile Thr Glu 305 310 315 320
- Phe Pro Asp Leu Thr Gly Thr Ala Asn Leu Glu Ser Leu Thr Leu Thr 325 330 335
- Gly Ala Gln Ile Ser Ser Leu Pro Gln Thr Val Cys Asn Gln Leu Pro 340 345 350
- Asn Leu Gln Val Leu Asp Leu Ser Tyr Asn Leu Leu Glu Asp Leu Pro 355 360 365
- Ser Phe Ser Val Cys Gln Lys Leu Gln Lys Ile Asp Leu Arq His Asn

	~ 7		_	~ -	- 1	-		_			~-		_	_	_	_
	Glu 385	Ile	Tyr	Glu	Ile	Lys 390	Va1	Asp	Thr	Phe	Gln 395		Leu	Leu	Ser	Let 400
	Arg	Ser	Leu	Asn	Leu 405	Ala	Trp	Asn	Lys	Ile 410	Ala	Ile	Ile	His	Pro 415	Asn
	Ala	Phe	Ser	Thr 420	Leu	Pro	Ser	Leu	Ile 425	_	Leu	Asp	Leu	Ser 430	Ser	Asn
	Leu	Leu	Ser 435	Ser	Phe	Pro	Ile	Thr 440	Gly	Leu	His	Gly	Leu 445	Thr	His	Leu
	Lys	Leu 450	Thr	Gly	Asn	His	Ala 455	Leu	Gln	Ser	Leu	Ile 460	Ser	Ser	Glu	Asn
	Phe 465	Pro	Glu	Leu	Lys	Val 470	Ile	Glu	Met	Pro	Tyr 475	Ala	Tyr	Gln	Cys	Cys 480
	Ala	Phe	Gly	Val	Cys 485	Glu	Asn	Ala	Tyr	Lys 490	Ile	Ser	Asn	Gln	Trp 495	Asn
	Lys	Gly	Asp	Asn 500	Ser	Ser	Met	Asp	Asp 505	Leu	His	Lys	Lys	Asp 510	Ala	Gly
	Met	Phe	Gln 515	Ala	Gln	Asp	Glu	Arg 520	Asp	Leu	Glu	Asp	Phe 525	Leu	Leu	Asp
	Phe	Glu 530	Glu	Asp	Leu	Lys	Ala 535	Leu	His	Ser	Val	Gln 540	Cys	Ser	Pro	Ser
	Pro 545	Gly	Pro	Phe	Lys	Pro 550	Cys	Glu	His	Leu	Leu 555	Asp	Gly	Trp	Leu	Ile 560
	Arg	Ile	Gly	Val	Trp 565	Thr	Ile	Ala	Val	Leu 570	Ala	Leu	Thr	Cys	Asn 575	Ala
	Leu	Val	Thr	Ser 580	Thr	Val	Phe	Arg	Ser 585	Pro	Leu	Tyr	Ile	Ser 590	Pro	Ile
	Lys	Leu	Leu 595	Ile	Gly	Val	Ile	Ala 600	Ala	Val	Asn	Met	Leu 605	Thr	Gly	Val
	Ser	Ser 610	Ala	Val	Leu	Ala	Gly 615	Val	Asp	Ala	Phe	Thr 620	Phe	Gly	Ser	Phe
	Ala 625	Arg	His	Gly	Ala	Trp 630	Trp	Glu	Asn	Gly	Val 635	Gly	Cys	His	Val	Ile 640
	Gly	Phe	Leu	Ser	Ile 645	Phe	Ala	Ser	Glu	Ser 650	Ser	Val	Phe	Leu	Leu 655	Thr
	Leu	Ala	Ala	Leu 660	Glu	Arg	Gly	Phe	Ser 665	Val	Lys	Tyr	Ser	Ala 670	Lys	Phe
(	Glu	Thr	Lys	Ala	Pro	Phe	Ser	Ser	Leu	Lys	Val	Ile	Ile	Leu	Leu	Cys

Ala Leu Leu Ala Leu Thr Met Ala Ala Val Pro Leu Leu Gly Gly Ser Lys Tyr Gly Ala Ser Pro Leu Cys Leu Pro Leu Pro Phe Gly Glu Pro 710 Ser Thr Met Gly Tyr Met Val Ala Leu Ile Leu Leu Asn Ser Leu Cys 725 730 Phe Leu Met Met Thr Ile Ala Tyr Thr Lys Leu Tyr Cys Asn Leu Asp 740 745 Lys Gly Asp Leu Glu Asn Ile Trp Asp Cys Ser Met Val Lys His Ile Ala Leu Leu Phe Thr Asn Cys Ile Leu Asn Cys Pro Val Ala Phe 775 Leu Ser Phe Ser Ser Leu Ile Asn Leu Thr Phe Ile Ser Pro Glu Val 790 Ile Lys Phe Ile Leu Leu Val Val Val Pro Leu Pro Ala Cys Leu Asn 805 810 Pro Leu Leu Tyr Ile Leu Phe Asn Pro His Phe Lys Glu Asp Leu Val 820 825 830 Ser Leu Arg Lys Gln Thr Tyr Val Trp Thr Arg Ser Lys His Pro Ser Leu Met Ser Ile Asn Ser Asp Asp Val Glu Lys Gln Ser Cys Asp Ser 855 Thr Gln Ala Leu Val Thr Phe Thr Ser Ser Ser Ile Thr Tyr Asp Leu 870 875 Pro Pro Ser Ser Val Pro Ser Pro Ala Tyr Pro Val Thr Glu Ser Cys 890

<210> 28

<211> 1531

<212> PRT

<213> Mus musculus

<400> 28

Met Ala Leu Thr Pro Gln Arg Gly Ser Ser Ser Gly Leu Ser Arg Pro 1 5 10 15

His Leu Ser Ser Val Ala Phe Val Pro Cys Leu

Glu Leu Trp Leu Leu Trp Ala Ala Trp Arg Leu Gly Ala Thr
20 25 30

Ala Cys Pro Ala Leu Cys Thr Cys Thr Gly Thr Thr Val Asp Cys His 40 Gly Thr Gly Leu Gln Ala Ile Pro Lys Asn Ile Pro Arg Asn Thr Glu 55 Arg Leu Glu Leu Asn Gly Asn Asn Ile Thr Arg Ile His Lys Asn Asp Phe Ala Gly Leu Lys Gln Leu Arg Val Leu Gln Leu Met Glu Asn Gln Ile Gly Ala Val Glu Arg Gly Ala Phe Asp Asp Met Lys Glu Leu Glu 105 Arg Leu Arg Leu Asn Arg Asn Gln Leu Gln Val Leu Pro Glu Leu Leu 120 Phe Gln Asn Asn Gln Ala Leu Ser Arg Leu Asp Leu Ser Glu Asn Phe 135 Leu Gln Ala Val Pro Arg Lys Ala Phe Arg Gly Ala Thr Asp Leu Lys 150 Asn Leu Gln Leu Asp Lys Asn Arg Ile Ser Cys Ile Glu Glu Gly Ala Phe Arg Ala Leu Arg Gly Leu Glu Val Leu Thr Leu Asn Asn Asn Asn 185 Ile Thr Thr Ile Pro Val Ser Ser Phe Asn His Met Pro Lys Leu Arg 200 Thr Phe Arg Leu His Ser Asn His Leu Phe Cys Asp Cys His Leu Ala Trp Leu Ser Gln Trp Leu Arg Gln Arg Pro Thr Ile Gly Leu Phe Thr 230 Gln Cys Ser Gly Pro Ala Ser Leu Arg Gly Leu Asn Val Ala Glu Val 245 Gln Lys Gly Glu Phe Ser Cys Ser Gly Gln Gly Glu Ala Ala Gly Ala Pro Ala Cys Thr Leu Ser Ser Gly Ser Cys Pro Ala Met Cys Ser Cys Ser Ser Gly Ile Val Asp Cys Arg Gly Lys Gly Leu Thr Ala Ile Pro Ala Asn Leu Pro Glu Thr Met Thr Glu Ile Arg Leu Glu Leu Asn Gly 310 Ile Lys Ser Ile Pro Pro Gly Ala Phe Ser Pro Tyr Arg Lys Leu Arg 325

Arg Ile Asp Leu Ser Asn Asn Gln Ile Ala Glu Ile Ala Pro Asp Ala 340 Phe Gln Gly Leu Arg Ser Leu Asn Ser Leu Val Leu Tyr Gly Asn Lys 360 Ile Thr Asp Leu Pro Arg Gly Val Phe Gly Gly Leu Tyr Thr Leu Gln 375 Leu Leu Leu Leu Asn Ala Asn Lys Ile Asn Cys Ile Arg Pro Asp Ala 390 Phe Gln Asp Leu Gln Asn Leu Ser Leu Leu Ser Leu Tyr Asp Asn Lys Ile Gln Ser Leu Ala Lys Gly Thr Phe Thr Ser Leu Arg Ala Ile Gln 425 Thr Leu His Leu Ala Gln Asn Pro Phe Ile Cys Asp Cys Asn Leu Lys 435 440 Trp Leu Ala Asp Phe Leu Arg Thr Asn Pro Ile Glu Thr Thr Gly Ala 455 Arg Cys Ala Ser Pro Arg Arg Leu Ala Asn Lys Arg Ile Gly Gln Ile Lys Ser Lys Lys Phe Arg Cys Ser Ala Lys Glu Gln Tyr Phe Ile Pro 490 Gly Thr Glu Asp Tyr His Leu Asn Ser Glu Cys Thr Ser Asp Val Ala 505 Cys Pro His Lys Cys Arg Cys Glu Ala Ser Val Val Glu Cys Ser Ser 520 Leu Lys Leu Ser Lys Ile Pro Glu Arg Ile Pro Gln Ser Thr Thr Glu 535 Leu Arg Leu Asn Asn Glu Ile Ser Ile Leu Glu Ala Thr Gly Leu 545 550 Phe Lys Lys Leu Ser His Leu Lys Lys Ile Asn Leu Ser Asn Asn Lys Val Ser Glu Ile Glu Asp Gly Thr Phe Glu Gly Ala Ala Ser Val Ser Glu Leu His Leu Thr Ala Asn Gln Leu Glu Ser Ile Arg Ser Gly Met Phe Arg Gly Leu Asp Gly Leu Arg Thr Leu Met Leu Arg Asn Asn Arg 615 Ile Ser Cys Ile His Asn Asp Ser Phe Thr Gly Leu Arg Asn Val Arg 625 630 635

Leu Leu Ser Leu Tyr Asp Asn His Ile Thr Thr Ile Ser Pro Gly Ala 645 Phe Asp Thr Leu Gln Ala Leu Ser Thr Leu Asn Leu Leu Ala Asn Pro 660 665 Phe Asn Cys Asn Cys His Leu Ser Trp Leu Gly Asp Trp Leu Arg Lys 680 Arg Lys Ile Val Thr Gly Asn Pro Arg Cys Gln Asn Pro Asp Phe Leu Arg Gln Ile Pro Leu Gln Asp Val Ala Phe Pro Asp Phe Arg Cys Glu 710 Glu Gly Gln Glu Val Gly Cys Leu Pro Arg Pro Gln Cys Pro Gln 730 Glu Cys Ala Cys Leu Asp Thr Val Val Arg Cys Ser Asn Lys His Leu 745 Gln Ala Leu Pro Lys Gly Ile Pro Lys Asn Val Thr Glu Leu Tyr Leu 760 Asp Gly Asn Gln Phe Thr Leu Val Pro Gly Gln Leu Ser Thr Phe Lys 770 775 Tyr Leu Gln Leu Val Asp Leu Ser Asn Asn Lys Ile Ser Ser Leu Ser 790 795 Asn Ser Ser Phe Thr Asn Met Ser Gln Leu Thr Thr Leu Ile Leu Ser 810 Tyr Asn Ala Leu Gln Cys Ile Pro Pro Leu Ala Phe Gln Arg Leu Arg Ser Leu Arg Leu Leu Ser Leu His Gly Asn Asp Val Ser Thr Leu Gln 840 Glu Gly Ile Phe Ala Asp Val Thr Ser Leu Ser His Leu Ala Ile Gly 850 855 Ala Asn Pro Leu Tyr Cys Asp Cys Arg Leu Arg Trp Leu Ser Ser Trp Val Lys Thr Gly Tyr Lys Glu Pro Gly Ile Ala Arg Cys Ala Gly Pro 885 Pro Glu Met Glu Gly Lys Leu Leu Thr Thr Pro Ala Lys Lys Phe 900 Glu Cys Gln Gly Pro Pro Ser Leu Ala Val Gln Ala Lys Cys Asp Pro 920 Cys Leu Ser Ser Pro Cys Gln Asn Gln Gly Thr Cys His Asn Asp Pro 930 935 940

- Leu Glu Val Tyr Arg Cys Thr Cys Pro Ser Gly Tyr Lys Gly Arg His 945 950 955 960
- Cys Glu Val Ser Leu Asp Gly Cys Ser Ser Asn Pro Cys Gly Asn Gly 965 970 975
- Gly Thr Cys His Ala Gln Glu Gly Glu Asp Ala Gly Phe Thr Cys Ser 980 985 990
- Cys Pro Ser Gly Phe Glu Gly Pro Thr Cys Gly Val Asp Thr Asp Asp 995 1000 1005
- Cys Val Lys His Ala Cys Val Asn Gly Gly Val Cys Val Asp Gly Val
  1010 1015 1020
- Gly Asn Tyr Thr Cys Gln Cys Pro Leu Gln Tyr Thr Gly Arg Ala Cys 1025 1030 1035 1040
- Glu Gln Leu Val Asp Phe Cys Ser Pro Asp Met Asn Pro Cys Gln His 1045 1050 1055
- Glu Ala Gln Cys Val Gly Thr Pro Asp Gly Pro Arg Cys Glu Cys Met 1060 1065 1070
- Leu Gly Tyr Thr Gly Asp Asn Cys Ser Glu Asn Gln Asp Asp Cys Lys 1075 1080 1085
- Asp His Lys Cys Gln Asn Gly Ala Gln Cys Val Asp Glu Val Asn Ser 1090 1095 1100
- Tyr Ala Cys Leu Cys Val Glu Gly Tyr Ser Gly Gln Leu Cys Glu Ile 1105 1110 1115 1120
- Pro Pro Ala Pro Arg Ser Ser Cys Glu Gly Thr Glu Cys Gln Asn Gly
  1125 1130 1135
- Ala Asn Cys Val Asp Gln Gly Ser Arg Pro Val Cys Gln Cys Leu Pro 1140 1150
- Gly Phe Gly Gly Pro Glu Cys Glu Lys Leu Leu Ser Val Asn Phe Val 1155 1160 1165
- Asp Arg Asp Thr Tyr Leu Gln Phe Thr Asp Leu Gln Asn Trp Pro Arg 1170 1175 1180
- Ala Asn Ile Thr Leu Gln Val Ser Thr Ala Glu Asp Asn Gly Ile Leu 1185 1190 1195 1200
- Leu Tyr Asn Gly Asp Asn Asp His Ile Ala Val Glu Leu Tyr Gln Gly
  1205 1210 1215
- His Val Arg Val Ser Tyr Asp Pro Gly Ser Tyr Pro Ser Ser Ala Ile 1220 1225 1230
- Tyr Ser Ala Glu Thr Ile Asn Asp Gly Gln Phe His Thr Val Glu Leu 1235 1240 1245

- Val Thr Phe Asp Gln Met Val Asn Leu Ser Ile Asp Gly Gly Ser Pro 1250 1255 1260
- Met Thr Met Asp Asn Phe Gly Lys His Tyr Thr Leu Asn Ser Glu Ala 1265 1270 1275 1280
- Pro Leu Tyr Val Gly Gly Met Pro Val Asp Val Asn Ser Ala Ala Phe 1285 1290 1295
- Arg Leu Trp Gln Ile Leu Asn Gly Thr Ser Phe His Gly Cys Ile Arg 1300 1305 1310
- Asn Leu Tyr Ile Asn Asn Glu Leu Gln Asp Phe Thr Lys Thr Gln Met 1315 1320 1325
- Lys Pro Gly Val Val Pro Gly Cys Glu Pro Cys Arg Lys Leu Tyr Cys 1330 1335 1340
- Leu His Gly Ile Cys Gln Pro Asn Ala Thr Pro Gly Pro Val Cys His 1345 1350 1355 1360
- Cys Glu Ala Gly Trp Gly Gly Leu His Cys Asp Gln Pro Val Asp Gly
  1365 1370 1375
- Pro Cys His Gly His Lys Cys Val His Gly Lys Cys Val Pro Leu Asp 1380 1385 1390
- Ala Leu Ala Tyr Ser Cys Gln Cys Gln Asp Gly Tyr Ser Gly Ala Leu 1395 1400 1405
- Cys Asn Gln Val Gly Ala Val Ala Glu Pro Cys Gly Gly Leu Gln Cys 1410 1420
- Leu His Gly His Cys Gln Ala Ser Ala Thr Lys Gly Ala His Cys Val 1425 1430 1435 1440
- Cys Ser Pro Gly Phe Ser Gly Glu Leu Cys Glu Gln Glu Ser Glu Cys 1445 1450 1455
- Arg Gly Asp Pro Val Arg Asp Phe His Arg Val Gln Arg Gly Tyr Ala 1460 1465 1470
- Ile Cys Gln Thr Thr Arg Pro Leu Ser Trp Val Glu Cys Arg Gly Ala 1475 1480 1485
- Cys Pro Gly Gln Gly Cys Cys Gln Gly Leu Arg Leu Lys Arg Arg Lys 1490 1495 1500
- Leu Thr Phe Glu Cys Ser Asp Gly Thr Ser Phe Ala Glu Glu Val Glu 1505 1510 1515 1520
- Lys Pro Thr Lys Cys Gly Cys Ala Gln Cys Val 1525 1530

<210> 29

<211> 1523

- <212> PRT
- <213> Drosophila melanogaster
- <400> 29
- Met Ala Pro Gly Trp Ala Gly Val Gly Ala Ala Val Arg Ala Arg Leu
  1 5 10 15
- Ala Leu Ala Leu Ala Ser Val Leu Ser Gly Pro Pro Ala Val 20 25 30
- Ala Cys Pro Thr Lys Cys Thr Cys Ser Ala Ala Ser Val Asp Cys His 35 40 45
- Gly Leu Gly Leu Arg Ala Val Pro Arg Gly Ile Pro Arg Asn Ala Glu
  50 55 60
- Arg Leu Asp Leu Asp Arg Asn Asn Ile Thr Arg Ile Thr Lys Met Asp 65 70 75 80
- Phe Ala Gly Leu Lys Asn Leu Arg Val Leu His Leu Glu Asp Asn Gln 85 90 95
- Val Ser Val Ile Glu Arg Gly Ala Phe Gln Asp Leu Lys Gln Leu Glu
  100 105 110
- Arg Leu Arg Leu Asn Lys Asn Lys Leu Gln Val Leu Pro Glu Leu Leu 115 120 125
- Phe Gln Ser Thr Pro Lys Leu Thr Arg Leu Asp Leu Ser Glu Asn Gln 130 135 140
- Ile Gln Gly Ile Pro Arg Lys Ala Phe Arg Gly Ile Thr Asp Val Lys
  145 150 155 160
- Asn Leu Gln Leu Asp Asn Asn His Ile Ser Cys Ile Glu Asp Gly Ala 165 170 175
- Phe Arg Ala Leu Arg Asp Leu Glu Ile Leu Thr Leu Asn Asn Asn Asn 180 185 190
- Ile Ser Arg Ile Leu Val Thr Ser Phe Asn His Met Pro Lys Ile Arg 195 200 205
- Thr Leu Arg Leu His Ser Asn His Leu Tyr Cys Asp Cys His Leu Ala 210 215 220
- Trp Leu Ser Asp Trp Leu Arg Gln Arg Arg Thr Val Gly Gln Phe Thr 225 230 235 240
- Leu Cys Met Ala Pro Val His Leu Arg Gly Phe Asn Val Ala Asp Val
  245 250 255
- Gln Lys Lys Glu Tyr Val Cys Pro Ala Pro His Ser Glu Pro Pro Ser 260 265 270
- Cys Asn Ala Asn Ser Ile Ser Cys Pro Ser Pro Cys Thr Cys Ser Asn 275 280 285

Asn Ile Val Asp Cys Arg Gly Lys Gly Leu Met Glu Ile Pro Ala Asn Leu Pro Glu Gly Ile Val Glu Ile Arg Leu Glu Gln Asn Ser Ile Lys 310 315 Ala Ile Pro Ala Gly Ala Phe Thr Gln Tyr Lys Lys Leu Lys Arg Ile Asp Ile Ser Lys Asn Gln Ile Ser Asp Ile Ala Pro Asp Ala Phe Gln 345 Gly Leu Lys Ser Leu Thr Ser Leu Val Leu Tyr Gly Asn Lys Ile Thr Glu Ile Ala Lys Gly Leu Phe Asp Gly Leu Val Ser Leu Gln Leu Leu Leu Leu Asn Ala Asn Lys Ile Asn Cys Leu Arg Val Asn Thr Phe Gln 390 395 Asp Leu Gln Asn Leu Asn Leu Leu Ser Leu Tyr Asp Asn Lys Leu Gln 405 Thr Ile Ser Lys Gly Leu Phe Ala Pro Leu Gln Ser Ile Gln Thr Leu 425 His Leu Ala Gln Asn Pro Phe Val Cys Asp Cys His Leu Lys Trp Leu 440 Ala Asp Tyr Leu Gln Asp Asn Pro Ile Glu Thr Ser Gly Ala Arg Cys Ser Ser Pro Arg Arg Leu Ala Asn Lys Arg Ile Ser Gln Ile Lys Ser 470 Lys Lys Phe Arg Cys Ser Gly Ser Glu Asp Tyr Arg Ser Arg Phe Ser 485 Ser Glu Cys Phe Met Asp Leu Val Cys Pro Glu Lys Cys Arg Cys Glu Gly Thr Ile Val Asp Cys Ser Asn Gln Lys Leu Val Arg Ile Pro Ser 515 520 525 His Leu Pro Glu Tyr Val Thr Asp Leu Arg Leu Asn Asp Asn Glu Val 530 Ser Val Leu Glu Ala Thr Gly Ile Phe Lys Lys Leu Pro Asn Leu Arg 550 555 Lys Ile Asn Leu Ser Asn Asn Lys Ile Lys Glu Val Arg Glu Gly Ala Phe Asp Gly Ala Ala Ser Val Gln Glu Leu Met Leu Thr Gly Asn Gln 580 585 590

Leu Glu Thr Val His Gly Arg Val Phe Arg Gly Leu Ser Gly Leu Lys 600 Thr Leu Met Leu Arg Ser Asn Leu Ile Gly Cys Val Ser Asn Asp Thr 615 Phe Ala Gly Leu Ser Ser Val Arg Leu Leu Ser Leu Tyr Asp Asn Arg 630 Ile Thr Thr Ile Thr Pro Gly Ala Phe Thr Thr Leu Val Ser Leu Ser 650 Thr Ile Asn Leu Leu Ser Asn Pro Phe Asn Cys Asn Cys His Leu Ala Trp Leu Gly Lys Trp Leu Arg Lys Arg Arg Ile Val Ser Gly Asn Pro Arg Cys Gln Lys Pro Phe Phe Leu Lys Glu Ile Pro Ile Gln Asp Val 695 Ala Ile Gln Asp Phe Thr Cys Asp Gly Asn Glu Glu Ser Ser Cys Gln 710 Leu Ser Pro Arg Cys Pro Glu Gln Cys Thr Cys Met Glu Thr Val Val Arg Cys Ser Asn Lys Gly Leu Arg Ala Leu Pro Arg Gly Met Pro Lys 740 745 Asp Val Thr Glu Leu Tyr Leu Glu Gly Asn His Leu Thr Ala Val Pro Arg Glu Leu Ser Ala Leu Arg His Leu Thr Leu Ile Asp Leu Ser Asn 775 Asn Ser Ile Ser Met Leu Thr Asn Tyr Thr Phe Ser Asn Met Ser His 790 Leu Ser Thr Leu Ile Leu Ser Tyr Asn Arg Leu Arg Cys Ile Pro Val His Ala Phe Asn Gly Leu Arg Ser Leu Arg Val Leu Thr Leu His Gly 820 825 Asn Asp Ile Ser Ser Val Pro Glu Gly Ser Phe Asn Asp Leu Thr Ser 835 Leu Ser His Leu Ala Leu Gly Thr Asn Pro Leu His Cys Asp Cys Ser 855 Leu Arg Trp Leu Ser Glu Trp Val Lys Ala Gly Tyr Lys Glu Pro Gly Ile Ala Arg Cys Ser Ser Pro Glu Pro Met Ala Asp Arg Leu Leu 885 890

- Thr Thr Pro Thr His Arg Phe Gln Cys Lys Gly Pro Val Asp Ile Asn 900 905 910
- Ile Val Ala Lys Cys Asn Ala Cys Leu Ser Ser Pro Cys Lys Asn Asn 915 920 925
- Gly Thr Cys Thr Gln Asp Pro Val Glu Leu Tyr Arg Cys Ala Cys Pro 930 935 940
- Tyr Ser Tyr Lys Gly Lys Asp Cys Thr Val Pro Ile Asn Thr Cys Ile 945 950 955 960
- Gln Asn Pro Cys Gln His Gly Gly Thr Cys His Leu Ser Asp Ser His 965 970 975
- Lys Asp Gly Phe Ser Cys Ser Cys Pro Leu Gly Phe Glu Gly Gln Arg 980 985 990
- Cys Glu Ile Asn Pro Asp Asp Cys Glu Asp Asn Asp Cys Glu Asn Asn 995 1000 1005
- Ala Thr Cys Val Asp Gly Ile Asn Asn Tyr Val Cys Ile Cys Pro Pro 1010 1015 1020
- Asn Tyr Thr Gly Glu Leu Cys Asp Glu Val Ile Asp His Cys Val Pro 1025 1030 1035 1040
- Glu Leu Asn Leu Cys Gln His Glu Ala Lys Cys Ile Pro Leu Asp Lys 1045 1050 1055
- Gly Phe Ser Cys Glu Cys Val Pro Gly Tyr Ser Gly Lys Leu Cys Glu 1060 1065 1070
- Thr Asp Asn Asp Asp Cys Val Ala His Lys Cys Arg His Gly Ala Gln 1075 1080 1085
- Cys Val Asp Thr Ile Asn Gly Tyr Thr Cys Thr Cys Pro Gln Gly Phe 1090 1095 1100
- Ser Gly Pro Phe Cys Glu His Pro Pro Pro Met Val Leu Leu Gln Thr 1105 1110 1115 1120
- Ser Pro Cys Asp Gln Tyr Glu Cys Gln Asn Gly Ala Gln Cys Ile Val 1125 1130 1135
- Val Gln Gln Glu Pro Thr Cys Arg Cys Pro Pro Gly Phe Ala Gly Pro 1140 1145 1150
- Arg Cys Glu Lys Leu Ile Thr Val Asn Phe Val Gly Lys Asp Ser Tyr 1155 1160 1165
- Val Glu Leu Ala Ser Ala Lys Val Arg Pro Gln Ala Asn Ile Ser Leu 1170 1175 1180
- Gln Val Ala Thr Asp Lys Asp Asn Gly Ile Leu Leu Tyr Lys Gly Asp 1185 1190 1195 1200

- Asn Asp Pro Leu Ala Leu Glu Leu Tyr Gln Gly His Val Arg Leu Val 1205 1210 1215
- Tyr Asp Ser Leu Ser Ser Pro Pro Thr Thr Val Tyr Ser Val Glu Thr
  1220 1225 1230
- Val Asn Asp Gly Gln Phe His Ser Val Glu Leu Val Thr Leu Asn Gln 1235 1240 1245
- Thr Leu Asn Leu Val Val Asp Lys Gly Thr Pro Lys Ser Leu Gly Lys 1250 1255 1260
- Leu Gln Lys Gln Pro Ala Val Gly Ile Asn Ser Pro Leu Tyr Leu Gly 1265 1270 1275 1280
- Gly Ile Pro Thr Ser Thr Gly Leu Ser Ala Leu Arg Gln Gly Thr Asp 1285 1290 1295
- Arg Pro Leu Gly Gly Phe His Gly Cys Ile His Glu Val Arg Ile Asn 1300 1305 1310
- Asn Glu Leu Gln Asp Phe Lys Ala Leu Pro Pro Gln Ser Leu Gly Val 1315 1320 1325
- Ser Pro Gly Cys Lys Ser Cys Thr Val Cys Lys His Gly Leu Cys Arg 1330 1335 1340
- Ser Val Glu Lys Asp Ser Val Val Cys Glu Cys Arg Pro Gly Trp Thr 1345 1350 1355 1360
- Gly Pro Leu Cys Asp Gln Glu Ala Arg Asp Pro Cys Leu Gly His Arg 1365 1370 1375
- Cys His His Gly Lys Cys Val Ala Thr Gly Thr Ser Tyr Met Cys Lys 1380 1385 1390
- Cys Ala Glu Gly Tyr Gly Gly Asp Leu Cys Asp Asn Lys Asn Asp Ser 1395 1400 1405
- Ala Asn Ala Cys Ser Ala Phe Lys Cys His His Gly Gln Cys His Ile 1410 1415 1420
- Ser Asp Gln Gly Glu Pro Tyr Cys Leu Cys Gln Pro Gly Phe Ser Gly 1425 1430 1435 1440
- Glu His Cys Gln Gln Glu Asn Pro Cys Leu Gly Gln Val Val Arg Glu 1445 1450 1455
- Val Ile Arg Arg Gln Lys Gly Tyr Ala Ser Cys Ala Thr Ala Ser Lys 1460 1465 1470
- Val Pro Ile Met Glu Cys Arg Gly Gly Cys Gly Pro Gln Cys Cys Gln
  1475 1480 1485
- Pro Thr Arg Ser Lys Arg Arg Lys Tyr Val Phe Gln Cys Thr Asp Gly 1490 1495 1500

Ser Ser Phe Val Glu Glu Val Glu Arg His Leu Glu Cys Gly Cys Leu 1505 1510 1515 1520

Ala Cys Ser

<210> 30

<211> 1534

<212> PRT

<213> Drosophila melanogaster

<400> 30

Met Ala Leu Thr Pro Gly Trp Gly Ser Ser Ala Gly Pro Val Arg Pro
1 5 10 15

Glu Leu Trp Leu Leu Trp Ala Ala Ala Trp Arg Leu Gly Ala Ser 20 25 30

Ala Cys Pro Ala Leu Cys Thr Cys Thr Gly Thr Thr Val Asp Cys His
35 40 45

Gly Thr Gly Leu Gln Ala Ile Pro Lys Asn Ile Pro Arg Asn Thr Glu
50 55 60

Arg Leu Glu Leu Asn Gly Asn Asn Ile Thr Arg Ile His Lys Asn Asp 65 70 75 80

Phe Ala Gly Leu Lys Gln Leu Arg Val Leu Gln Leu Met Glu Asn Gln
85 90 95

Ile Gly Ala Val Glu Arg Gly Ala Phe Asp Asp Met Lys Glu Leu Glu
100 105 110

Arg Leu Arg Leu Asn Arg Asn Gln Leu His Met Leu Pro Glu Leu Leu 115 120 125

Phe Gln Asn Asn Gln Ala Leu Ser Arg Leu Asp Leu Ser Glu Asn Ala 130 135 140

Ile Gln Ala Ile Pro Arg Lys Ala Phe Arg Gly Ala Thr Asp Leu Lys 145 150 155 160

Asn Leu Arg Leu Asp Lys Asn Gln Ile Ser Cys Ile Glu Glu Gly Ala 165 170 175

Phe Arg Ala Leu Arg Gly Leu Glu Val Leu Thr Leu Asn Asn Asn Asn 180

Ile Thr Thr Ile Pro Val Ser Ser Phe Asn His Met Pro Lys Leu Arg 195 200 205

Thr Phe Arg Leu His Ser Asn His Leu Phe Cys Asp Cys His Leu Ala 210 215 220

Trp Leu Ser Gln Trp Leu Arg Gln Arg Pro Thr Ile Gly Leu Phe Thr

225	230	235	240

- Gln Cys Ser Gly Pro Ala Ser Leu Arg Gly Leu Asn Val Ala Glu Val 245 250 255
- Gln Lys Ser Glu Phe Ser Cys Ser Gly Gln Gly Glu Ala Gly Arg Val 260 265 270
- Pro Thr Cys Thr Leu Ser Ser Gly Ser Cys Pro Ala Met Cys Thr Cys 275 280 285
- Ser Asn Gly Ile Val Asp Cys Arg Gly Lys Gly Leu Thr Ala Ile Pro 290 295 300
- Ala Asn Leu Pro Glu Thr Met Thr Glu Ile Arg Leu Glu Leu Asn Gly 305 310 315 320
- Ile Lys Ser Ile Pro Pro Gly Ala Phe Ser Pro Tyr Arg Lys Leu Arg 325 330 335
- Arg Ile Asp Leu Ser Asn Asn Gln Ile Ala Glu Ile Ala Pro Asp Ala 340 345 350
- Phe Gln Gly Leu Arg Ser Leu Asn Ser Leu Val Leu Tyr Gly Asn Lys 355 360 365
- Ile Thr Asp Leu Pro Arg Gly Val Phe Gly Gly Leu Tyr Thr Leu Gln 370 375 380
- Leu Leu Leu Leu Asn Ala Asn Lys Ile Asn Cys Ile Arg Pro Asp Ala 385 390 395 400
- Phe Gln Asp Leu Gln Asn Leu Ser Leu Leu Ser Leu Tyr Asp Asn Lys 405 410 415
- Ile Gln Ser Leu Ala Lys Gly Thr Phe Thr Ser Leu Arg Ala Ile Gln 420 425 430
- Thr Leu His Leu Ala Gln Asn Pro Phe Ile Cys Asp Cys Asn Leu Lys
  435 440 445
- Trp Leu Ala Asp Phe Leu Arg Thr Asn Pro Ile Glu Thr Ser Gly Ala 450 455 460
- Arg Cys Ala Ser Pro Arg Arg Leu Ala Asn Lys Arg Ile Gly Gln Ile 465 470 475 480
- Lys Ser Lys Lys Phe Arg Cys Ser Ala Lys Glu Gln Tyr Phe Ile Pro 485 490 495
- Gly Thr Glu Asp Tyr Gln Leu Asn Ser Glu Cys Asn Ser Asp Val Val 500 510
- Cys Pro His Lys Cys Arg Cys Glu Ala Asn Val Val Glu Cys Ser Ser 515 520 525
- Leu Lys Leu Thr Lys Ile Pro Glu Arg Ile Pro Gln Ser Thr Ala Glu

Leu Arg Leu Asn Asn Glu Ile Ser Ile Leu Glu Ala Thr Gly Met 550 555 Phe Lys Lys Leu Thr His Leu Lys Lys Ile Asn Leu Ser Asn Asn Lys Val Ser Glu Ile Glu Asp Gly Ala Phe Glu Gly Ala Ala Ser Val Ser Glu Leu His Leu Thr Ala Asn Gln Leu Glu Ser Ile Arg Ser Gly Met 600 Phe Arg Gly Leu Asp Gly Leu Arg Thr Leu Met Leu Arg Asn Asn Arg 615 Ile Ser Cys Ile His Asn Asp Ser Phe Thr Gly Leu Arg Asn Val Arg 630 635 Leu Leu Ser Leu Tyr Asp Asn Gln Ile Thr Thr Val Ser Pro Gly Ala 650 Phe Asp Thr Leu Gln Ser Leu Ser Thr Leu Asn Leu Leu Ala Asn Pro Phe Asn Cys Asn Cys Gln Leu Ala Trp Leu Gly Gly Trp Leu Arg Lys 675 680 Arg Lys Ile Val Thr Gly Asn Pro Arg Cys Gln Asn Pro Asp Phe Leu Arg Gln Ile Pro Leu Gln Asp Val Ala Phe Pro Asp Phe Arg Cys Glu Glu Gly Gln Glu Gly Gly Cys Leu Pro Arg Pro Gln Cys Pro Gln 725 730 Glu Cys Ala Cys Leu Asp Thr Val Val Arg Cys Ser Asn Lys His Leu 745 Arg Ala Leu Pro Lys Gly Ile Pro Lys Asn Val Thr Glu Leu Tyr Leu 760 Asp Gly Asn Gln Phe Thr Leu Val Pro Gly Gln Leu Ser Thr Phe Lys Tyr Leu Gln Leu Val Asp Leu Ser Asn Asn Lys Ile Ser Ser Leu Ser 790 Asn Ser Ser Phe Thr Asn Met Ser Gln Leu Thr Thr Leu Ile Leu Ser 805 Tyr Asn Ala Leu Gln Cys Ile Pro Pro Leu Ala Phe Gln Gly Leu Arq 825

Ser Leu Arg Leu Leu Ser Leu His Gly Asn Asp Ile Ser Thr Leu Gln

- Glu Gly Ile Phe Ala Asp Val Thr Ser Leu Ser His Leu Ala Ile Gly 855 Ala Asn Pro Leu Tyr Cys Asp Cys His Leu Arg Trp Leu Ser Ser Trp Val Lys Thr Gly Tyr Lys Glu Pro Gly Ile Ala Arg Cys Ala Gly Pro Gln Asp Met Glu Gly Lys Leu Leu Leu Thr Thr Pro Ala Lys Lys Phe 900 905 Glu Cys Gln Gly Pro Pro Thr Leu Ala Val Gln Ala Lys Cys Asp Leu 920 Cys Leu Ser Ser Pro Cys Gln Asn Gln Gly Thr Cys His Asn Asp Pro 935 Leu Glu Val Tyr Arg Cys Ala Cys Pro Ser Gly Tyr Lys Gly Arg Asp Cys Glu Val Ser Leu Asn Ser Cys Ser Ser Gly Pro Cys Glu Asn Gly Gly Thr Cys His Ala Gln Glu Gly Glu Asp Ala Pro Phe Thr Cys Ser 980 985 Cys Pro Thr Gly Phe Glu Gly Pro Thr Cys Gly Val Asn Thr Asp Asp Cys Val Asp His Ala Cys Ala Asn Gly Gly Val Cys Val Asp Gly Val 1015 Gly Asn Tyr Thr Cys Gln Cys Pro Leu Gln Tyr Glu Gly Lys Ala Cys 1025 1030 1035 Glu Gln Leu Val Asp Leu Cys Ser Pro Asp Leu Asn Pro Cys Gln His 1045 1050 Glu Ala Gln Cys Val Gly Thr Pro Asp Gly Pro Arg Cys Glu Cys Met 1065 Pro Gly Tyr Ala Gly Asp Asn Cys Ser Glu Asn Gln Asp Asp Cys Arq
- Asp His Arg Cys Gln Asn Gly Ala Gln Cys Met Asp Glu Val Asn Ser 1090 1095 1100
- Tyr Ser Cys Leu Cys Ala Glu Gly Tyr Ser Gly Gln Leu Cys Glu Ile 1105 1110 1115 1120
- Pro Pro His Leu Pro Ala Pro Lys Ser Pro Cys Glu Gly Thr Glu Cys 1125 1130 1135
- Gln Asn Gly Ala Asn Cys Val Asp Gln Gly Asn Arg Pro Val Cys Gln

- Cys Leu Pro Gly Phe Gly Gly Pro Glu Cys Glu Lys Leu Leu Ser Val 1155 1160 1165
- Asn Phe Val Asp Arg Asp Thr Tyr Leu Gln Phe Thr Asp Leu Gln Asn 1170 1175 1180
- Trp Pro Arg Ala Asn Ile Thr Leu Gln Val Ser Thr Ala Glu Asp Asn 1185 1190 1195 1200
- Gly Ile Leu Leu Tyr Asn Gly Asp Asn Asp His Ile Ala Val Glu Leu 1205 1210 1215
- Tyr Gln Gly His Val Arg Val Ser Tyr Asp Pro Gly Ser Tyr Pro Ser 1220 1225 1230
- Ser Ala Ile Tyr Ser Ala Glu Thr Ile Asn Asp Gly Gln Phe His Thr 1235 1240 1245
- Val Glu Leu Val Ala Phe Asp Gln Met Val Asn Leu Ser Ile Asp Gly 1250 1255 1260
- Gly Ser Pro Met Thr Met Asp Asn Phe Gly Lys His Tyr Thr Leu Asn 1265 1270 1275 1280
- Ser Glu Ala Pro Leu Tyr Val Gly Gly Met Pro Val Asp Val Asn Ser 1285 1290 1295
- Ala Ala Phe Arg Leu Trp Gln Ile Leu Asn Gly Thr Gly Phe His Gly
  1300 1305 1310
- Cys Ile Arg Asn Leu Tyr Ile Asn Asn Glu Leu Gln Asp Phe Thr Lys 1315 1320 1325
- Thr Gln Met Lys Pro Gly Val Val Pro Gly Cys Glu Pro Cys Arg Lys 1330 1335 1340
- Leu Tyr Cys Leu His Gly Ile Cys Gln Pro Asn Ala Thr Pro Gly Pro 1345 1350 1355 1360
- Met Cys His Cys Glu Ala Gly Trp Val Gly Leu His Cys Asp Gln Pro 1365 1370 1375
- Ala Asp Gly Pro Cys His Gly His Lys Cys Val His Gly Gln Cys Val 1380 1385 1390
- Pro Leu Asp Ala Leu Ser Tyr Ser Cys Gln Cys Gln Asp Gly Tyr Ser 1395 1400 1405
- Gly Ala Leu Cys Asn Gln Ala Gly Ala Leu Ala Glu Pro Cys Arg Gly 1410 1415 1420
- Leu Gln Cys Leu His Gly His Cys Gln Ala Ser Gly Thr Lys Gly Ala 1425 1430 1435 1440
- His Cys Val Cys Asp Pro Gly Phe Ser Gly Glu Leu Cys Glu Gln Glu

Ser Glu Cys Arg Gly Asp Pro Val Arg Asp Phe His Gln Val Gln Arg 1460 1465 1470

Gly Tyr Ala Ile Cys Gln Thr Thr Arg Pro Leu Ser Trp Val Glu Cys 1475 1480 1485

Arg Gly Ser Cys Pro Gly Gln Gly Cys Cys Gln Gly Leu Arg Leu Lys 1490 1495 1500

Arg Arg Lys Phe Thr Phe Glu Cys Ser Asp Gly Thr Ser Phe Ala Glu 1505 1510 1515 1520

Glu Val Glu Lys Pro Thr Lys Cys Gly Cys Ala Leu Cys Ala 1525 1530

<210> 31

<211> 1531

<2:12> PRT

<213> Rattus norvegicus

<400> 31

Met Ala Leu Thr Pro Gln Arg Gly Ser Ser Ser Gly Leu Ser Arg Pro 1 5 10 15

Glu Leu Trp Leu Leu Trp Ala Ala Ala Trp Arg Leu Gly Ala Thr
20 25 30

Ala Cys Pro Ala Leu Cys Thr Cys Thr Gly Thr Thr Val Asp Cys His 35 40 45

Gly Thr Gly Leu Gln Ala Ile Pro Lys Asn Ile Pro Arg Asn Thr Glu

Arg Leu Glu Leu Asn Gly Asn Asn Ile Thr Arg Ile His Lys Asn Asp 65 70 75 80

Phe Ala Gly Leu Lys Gln Leu Arg Val Leu Gln Leu Met Glu Asn Gln
85 90 95

Ile Gly Ala Val Glu Arg Gly Ala Phe Asp Asp Met Lys Glu Leu Glu 100 105 110

Arg Leu Arg Leu Asn Arg Asn Gln Leu Gln Val Leu Pro Glu Leu Leu 115 120 125

Phe Gln Asn Asn Gln Ala Leu Ser Arg Leu Asp Leu Ser Glu Asn Ser 130 135 140

Leu Gln Ala Val Pro Arg Lys Ala Phe Arg Gly Ala Thr Asp Leu Lys 145 150 155 160

Asn Leu Gln Leu Asp Lys Asn Gln Ile Ser Cys Ile Glu Glu Gly Ala 165 170 175

Phe Arg Ala Leu Arg Gly Leu Glu Val Leu Thr Leu Asn Asn Asn Asn Ile Thr Thr Ile Pro Val Ser Ser Phe Asn His Met Pro Lys Leu Arg 200 Thr Phe Arg Leu His Ser Asn His Leu Phe Cys Asp Cys His Leu Ala 215 Trp Leu Ser Gln Trp Leu Arg Gln Arg Pro Thr Ile Gly Leu Phe Thr 225 230 235 Gln Cys Ser Gly Pro Ala Ser Leu Arg Gly Leu Asn Val Ala Glu Val Gln Lys Ser Glu Phe Ser Cys Ser Gly Gln Gly Glu Ala Ala Gln Val 265 Pro Ala Cys Thr Leu Ser Ser Gly Ser Cys Pro Ala Met Cys Ser Cys 275 Ser Asn Gly Ile Val Asp Cys Arg Gly Lys Gly Leu Thr Ala Ile Pro 295 Ala Asn Leu Pro Glu Thr Met Thr Glu Ile Arg Leu Glu Leu Asn Gly 305 310 315 Ile Lys Ser Ile Pro Pro Gly Ala Phe Ser Pro Tyr Arg Lys Leu Arg Arg Ile Asp Leu Ser Asn Asn Gln Ile Ala Glu Ile Ala Pro Asp Ala 340 Phe Gln Gly Leu Arg Ser Leu Asn Ser Leu Val Leu Tyr Gly Asn Lys 355 360 Ile Thr Asp Leu Pro Arg Gly Val Phe Gly Gly Leu Tyr Thr Leu Gln 375 Leu Leu Leu Asn Ala Asn Lys Ile Asn Cys Ile Arg Pro Asp Ala 385 400 Phe Gln Asp Leu Gln Asn Leu Ser Leu Ser Leu Tyr Asp Asn Lys 405 Ile Gln Ser Leu Ala Lys Gly Thr Phe Thr Ser Leu Arg Ala Ile Gln 425 Thr Leu His Leu Ala Gln Asn Pro Phe Ile Cys Asp Cys Asn Leu Lys 435 440 Trp Leu Ala Asp Phe Leu Arg Thr Asn Pro Ile Glu Thr Thr Gly Ala 455 Arg Cys Ala Ser Pro Arg Arg Leu Ala Asn Lys Arg Ile Gly Gln Ile 470 475 480

Lys Ser Lys Lys Phe Arg Cys Ser Ala Lys Glu Gln Tyr Phe Ile Pro Gly Thr Glu Asp Tyr His Leu Asn Ser Glu Cys Thr Ser Asp Val Ala 500 Cys Pro His Lys Cys Arg Cys Glu Ala Ser Val Val Glu Cys Ser Gly 520 Leu Lys Leu Ser Lys Ile Pro Glu Arg Ile Pro Gln Ser Thr Thr Glu 530 535 Leu Arg Leu Asn Asn Asn Glu Ile Ser Ile Leu Glu Ala Thr Gly Leu 550 Phe Lys Lys Leu Ser His Leu Lys Lys Ile Asn Leu Ser Asn Asn Lys 570 Val Ser Glu Ile Glu Asp Gly Thr Phe Glu Gly Ala Thr Ser Val Ser 580 Glu Leu His Leu Thr Ala Asn Gln Leu Glu Ser Val Arg Ser Gly Met 600 Phe Arg Gly Leu Asp Gly Leu Arg Thr Leu Met Leu Arg Asn Asn Arg 610 615 Ile Ser Cys Ile His Asn Asp Ser Phe Thr Gly Leu Arg Asn Val Arg Leu Leu Ser Leu Tyr Asp Asn His Ile Thr Thr Ile Ser Pro Gly Ala 650 Phe Asp Thr Leu Gln Ala Leu Ser Thr Leu Asn Leu Leu Ala Asn Pro 660 665 Phe Asn Cys Asn Cys Gln Leu Ala Trp Leu Gly Asp Trp Leu Arg Lys 680 Arg Lys Ile Val Thr Gly Asn Pro Arg Cys Gln Asn Pro Asp Phe Leu 690 Arg Gln Ile Pro Leu Gln Asp Val Ala Phe Pro Asp Phe Arg Cys Glu Glu Gly Gln Glu Val Gly Cys Leu Pro Arg Pro Gln Cys Pro Gln 730 Glu Cys Ala Cys Leu Asp Thr Val Val Arg Cys Ser Asn Lys His Leu 740 Gln Ala Leu Pro Lys Gly Ile Pro Lys Asn Val Thr Glu Leu Tyr Leu 760 Asp Gly Asn Gln Phe Thr Leu Val Pro Gly Gln Leu Ser Thr Phe Lys

775

770

Tyr Leu Gln Leu Val Asp Leu Ser Asn Asn Lys Ile Ser Ser Leu Ser 785 790 Asn Ser Ser Phe Thr Asn Met Ser Gln Leu Thr Thr Leu Ile Leu Ser 805 810 Tyr Asn Ala Leu Gln Cys Ile Pro Pro Leu Ala Phe Gln Gly Leu Arg 825 Ser Leu Arg Leu Leu Ser Leu His Gly Asn Asp Val Ser Thr Leu Gln 840 Glu Gly Ile Phe Ala Asp Val Thr Ser Leu Ser His Leu Ala Ile Gly 855 Ala Asn Pro Leu Tyr Cys Asp Cys His Leu Arg Trp Leu Ser Ser Trp 870 Val Lys Thr Gly Tyr Lys Glu Pro Gly Ile Ala Arg Cys Ala Gly Pro 885 890 Pro Glu Met Glu Gly Lys Leu Leu Thr Thr Pro Ala Lys Lys Phe 905 Glu Cys Gln Gly Pro Pro Ser Leu Ala Val Gln Ala Lys Cys Asp Pro Cys Leu Ser Ser Pro Cys Gln Asn Gln Gly Thr Cys His Asn Asp Pro 935 Leu Glu Val Tyr Arg Cys Thr Cys Pro Ser Gly Tyr Lys Gly Arg Asn 950 955 Cys Glu Val Ser Leu Asp Ser Cys Ser Ser Asn Pro Cys Gly Asn Gly 965 Gly Thr Cys His Ala Gln Glu Gly Glu Asp Ala Gly Phe Thr Cys Ser Cys Pro Ser Gly Phe Glu Gly Leu Thr Cys Gly Met Asn Thr Asp Asp 995 1000 Cys Val Lys His Asp Cys Val Asn Gly Gly Val Cys Val Asp Gly Ile 1015 Gly Asn Tyr Thr Cys Gln Cys Pro Leu Gln Tyr Thr Gly Arg Ala Cys 1030 1035 Glu Gln Leu Val Asp Phe Cys Ser Pro Asp Leu Asn Pro Cys Gln His 1045 1050 Glu Ala Gln Cys Val Gly Thr Pro Glu Gly Pro Arg Cys Glu Cys Val 1065 Pro Gly Tyr Thr Gly Asp Asn Cys Ser Lys Asn Gln Asp Asp Cys Lys

1080

1075

- Asp His Gln Cys Gln Asn Gly Ala Gln Cys Val Asp Glu Ile Asn Ser 1090 1095 1100
- Tyr Ala Cys Leu Cys Ala Glu Gly Tyr Ser Gly Gln Leu Cys Glu Ile 1105 1110 1115 1120
- Pro Pro Ala Pro Arg Asn Ser Cys Glu Gly Thr Glu Cys Gln Asn Gly
  1125 1130 1135
- Ala Asn Cys Val Asp Gln Gly Ser Arg Pro Val Cys Gln Cys Leu Pro 1140 1145 1150
- Gly Phe Gly Gly Pro Glu Cys Glu Lys Leu Leu Ser Val Asn Phe Val 1155 1160 1165
- Asp Arg Asp Thr Tyr Leu Gln Phe Thr Asp Leu Gln Asn Trp Pro Arg 1170 1175 1180
- Ala Asn Ile Thr Leu Gln Val Ser Thr Ala Glu Asp Asn Gly Ile Leu 1185 1190 1195 1200
- Leu Tyr Asn Gly Asp Asn Asp His Ile Ala Val Glu Leu Tyr Gln Gly
  1205 1210 1215
- His Val Arg Val Ser Tyr Asp Pro Gly Ser Tyr Pro Ser Ser Ala Ile 1220 1225 1230
- Tyr Ser Ala Glu Thr Ile Asn Asp Gly Gln Phe His Thr Val Glu Leu 1235 1240 1245
- Val Thr Phe Asp Gln Met Val Asn Leu Ser Ile Asp Gly Gly Ser Pro 1250 1255 1260
- Met Thr Met Asp Asn Phe Gly Lys His Tyr Thr Leu Asn Ser Glu Ala 1265 1270 1275 1280
- Pro Leu Tyr Val Gly Gly Met Pro Val Asp Val Asn Ser Ala Ala Phe 1285 1290 1295
- Arg Leu Trp Gln Ile Leu Asn Gly Thr Ser Phe His Gly Cys Ile Arg 1300 1305 1310
- Asn Leu Tyr Ile Asn Asn Glu Leu Gln Asp Phe Thr Lys Thr Gln Met 1315 1320 1325
- Lys Pro Gly Val Val Pro Gly Cys Glu Pro Cys Arg Lys Leu Tyr Cys 1330 1335 1340
- Leu His Gly Ile Cys Gln Pro Asn Ala Thr Pro Gly Pro Val Cys His 1345 1350 1355 1360
- Cys Glu Ala Gly Trp Gly Gly Leu His Cys Asp Gln Pro Val Asp Gly
  1365 1370 1375
- Pro Cys His Gly His Lys Cys Val His Gly Lys Cys Val Pro Leu Asp 1380 1385 1390

- Ala Leu Ala Tyr Ser Cys Gln Cys Gln Asp Gly Tyr Ser Gly Ala Leu 1395 1400 1405
- Cys Asn Gln Val Gly Ala Val Ala Glu Pro Cys Gly Gly Leu Gln Cys 1410 1415 1420
- Leu His Gly His Cys Gln Ala Ser Ala Thr Arg Gly Ala His Cys Val 1425 1430 1435 1440
- Cys Ser Pro Gly Phe Ser Gly Glu Leu Cys Glu Gln Glu Ser Glu Cys 1445 1450 1455
- Arg Gly Asp Pro Val Arg Asp Phe His Arg Val Gln Arg Gly Tyr Ala 1460 1465 1470
- Ile Cys Gln Thr Thr Arg Pro Leu Ser Trp Val Glu Cys Arg Gly Ala 1475 1480 1485
- Cys Pro Gly Gln Gly Cys Cys Gln Gly Leu Arg Leu Lys Arg Arg Lys 1490 1495 1500
- Leu Thr Phe Glu Cys Ser Asp Gly Thr Ser Phe Ala Glu Glu Val Glu 1505 1510 1515 1520
- Lys Pro Thr Lys Cys Gly Cys Ala Pro Cys Ala 1525 1530

<210> 32

<211> 1512

<212> PRT

<213> Danio rerio

<400> 32

- Met Phe Val Leu Lys Ser Val Val Leu Cys Ala Leu Leu Cys Gly Ala 1 5 10 15
- Gly Ala Gln Ser Cys Pro Ser Gln Cys Ser Cys Ser Gly Thr Ala Val 20 25 30
- Asp Cys His Gly Gln Ser Leu Arg Ser Val Pro Arg Asn Ile Pro Arg 35 40 45
- Asn Val Glu Arg Leu Asp Leu Asn Ala Asn Asn Leu Thr Lys Ile Thr 50 55 60
- Lys Ala Asp Phe Ala Gly Leu Lys Asn Leu Arg Val Leu Gln Leu Met 65 70 75 80
- Glu Asn Lys Ile Ser Ser Ile Glu Arg Gly Ala Phe Gln Asp Leu Gln
  85 90 95
- Glu Leu Glu Arg Leu Arg Leu Asn Arg Asn Asn Leu Gln Val Leu Pro 100 105 110
- Glu Leu Leu Phe Leu Gly Thr Thr Lys Leu Phe Arg Leu Asp Leu Ser 115 120 125

Glu	Asn 130	Gln	Ile	Gln	Gly	Ile 135	Pro	Arg	Lys	Ala	Phe 140	Arg	Gly	Ser	Thr
Glu 145	Ile	Lys	Asn	Leu	Gln 150	Leu	Asp	Tyr	Asn	Gln 155	Ile	Ser	Cys	Ile	Glu 160
Asp	Gly	Ala	Phe	Arg 165	Ala	Leu	Gly	Asp	Leu 170		Val	Leu	Thr	Leu 175	Asn
Asn	Asn	Asn	Ile 180	Ser	Arg	Leu	Ser	Val 185	Ala	Ser	Phe	Asn	His 190	Met	Pro
Lys	Leu	Arg 195	Thr	Phe	Arg	Leu	His 200	Ser	Asn	Asn	Leu	Leu 205	Cys	Asp	Cys
Asn	Val 210	Ala	Trp	Leu	Ser	Asp 215	Trp	Leu	Arg	Gln	Arg 220	Pro	Arg	Leu	Gly
Leu 225	Tyr	Thr	Gln	Cys	Met 230	Ala	Pro	Pro	Ser	Leu 235	Arg	Gly	His	Asn	Ile 240
Ala	Glu	Val	Gln	Lys 245	Lys	Glu	Phe	Met	Cys 250	Thr	Gly	Pro	Gln	Ser 255	His
Ser	Ser	Cys	Ser 260	Val	Leu	Gln	Cys	Pro 265	Glu	Leu	Cys	Thr	Cys 270	Ser	Asn
Asn	Val	Val 275	Asp	Cys	Arg	Gly	Lys 280	Gly	Leu	Thr	Glu	Ile 285	Pro	Thr	Asn
Leu	Pro 290	Glu	Thr	Ile	Thr	Glu 295	Ile	Arg	Leu	Glu	Gln 300	Asn	Ser	Ile	Lys
Ile 305	Ile	Pro	Ala	Gly	Ala 310	Phe	Ala	Pro	Tyr	Lys 315	Arg	Leu	Arg	Arg	Ile 320
Asp	Leu	Ser	Asn	Asn 325	Gln	Ile	Thr	Glu	Leu 330	Ala	Ser	Asp	Ser	Phe 335	Gln
Gly	Leu	Arg	Ser 340	Leu	Asn	Ser	Leu	Val 345	Leu	Tyr	Gly	Asn	Lys 350	Ile	Thr
Glu	Leu	Pro 355	rys	Gly	Leu	Phe	Asp 360	Gly	Leu	Phe	Ser	Leu 365	Gln	Leu	Leu
Leu	Leu 370	Asn	Ala	Asn	Lys	Ile 375	Asn	Cys	Leu	Arg	Val 380	Asp	Ser	Phe	Gln
Asp 385	Leu	Gln	Asn	Leu	Asn 390	Leu	Leu	Ser	Leu	Tyr 395	Asp	Asn	Lys	Leu	Gln 400
Thr	Ile	Ala	Lys	Gly 405	Thr	Phe	Ser	Ser	Leu 410	Arg	Ala	Ile	Gln	Thr 415	Leu
His	Leu	Ala	Gln 420	Asn	Pro	Phe	Met	Cys 425	Asp	Cys	His	Leu	Lys 430	Trp	Leu

Ala Asp Tyr Leu Gln Asp Asn Pro Ile Glu Thr Ser Gly Ala Arg Cys 440 Thr Ser Pro Arg Arg Leu Ala Asn Lys Arg Ile Gly Gln Ile Lys Ser 455 Lys Lys Phe Arg Cys Ser Gly Val Glu Asp Tyr Arg Ser Lys Leu Gly Gly Asp Cys Phe Ala Asp Leu Ala Cys Pro Glu Lys Cys Arg Cys Glu 490 Gly Thr Thr Val Asp Cys Ser Gly Gln Lys Leu Thr Lys Ile Pro Asp 505 His Ile Pro Gln Tyr Thr Ala Glu Leu Arg Leu Asn Asn Asn Glu Phe 520 Thr Val Leu Glu Ala Thr Gly Ile Phe Lys Lys Leu Pro Gln Leu Arg 535 Lys Ile Asn Leu Ser Asn Asn Lys Ile Thr Asp Ile Glu Glu Gly Thr 545 550 Phe Glu Gly Ala Ser Gly Val Asn Glu Leu Ile Leu Thr Ser Asn Arg 570 Leu Glu Gly Val His Tyr Ser Met Leu Lys Gly Leu Gly Gly Leu Arg 585 Thr Leu Met Leu Arg Ser Asn Arg Ile Ser Cys Val Asn Asn Gly Ser Phe Thr Gly Leu Ser Ser Val Arg Leu Leu Ser Leu Tyr Asp Asn Leu 615 Ile Thr Ser Met Ser Pro Gly Ala Phe Asp Thr Leu His Ser Leu Ser 625 630 Thr Leu Asn Leu Leu Ala Asn Pro Phe Asn Cys Asn Cys His Leu Ala 650 Trp Leu Gly Glu Trp Leu Arg Lys Lys Arg Ile Val Thr Gly Asn Pro 660 Arg Cys Gln Ser Pro Tyr Phe Leu Lys Glu Ile Pro Ile Gln Asp Val Ala Val Gln Asp Phe Ala Cys Glu Glu Gly Asn Asp Glu Asn Ser Cys 695 Ser Pro Leu Ala Arg Cys Pro Ala Glu Cys Ser Cys Leu Asp Thr Val 705 710 Val Arg Cys Ser Asn Lys Gly Leu Lys Val Leu Pro Lys Gly Ile Pro 725 730

Arg Asp Val Thr Glu Leu Tyr Leu Asp Gly Asn Glu Phe Thr Gln Val 745 Pro Leu Glu Leu Ser Asn Tyr Lys His Leu Thr Leu Ile Asp Leu Ser 760 Asn Asn Gln Ile Ser Thr Leu Ser Asn His Ser Phe Ser Asn Met Ser Glu Leu Leu Thr Leu Ile Leu Ser Tyr Asn Arg Leu Arg Cys Ile Pro 790 Ala Lys Ala Phe Asp Gly Leu Lys Ser Leu Arg Leu Leu Ser Leu His 805 810 Gly Asn Asp Ile Ala Val Ile Pro Asp Gly Ala Phe Lys Asp Leu Ser 820 825 Ser Leu Ser His Leu Ala Leu Gly Ala Asn Pro Leu Tyr Cys Asp Cys 840 His Met Gln Trp Leu Ser Asp Trp Val Lys Ser Gly Tyr Lys Glu Pro 850 Gly Ile Ala Arg Cys Thr Gly Pro Gly Asp Met Thr Asp Lys Leu Leu Leu Thr Thr Pro Ser Lys Lys Phe Thr Cys Thr Gly Pro Val Asp Val 890 Ser Ile Leu Ala Lys Cys Asn Pro Cys Leu Ser Asn Pro Cys Lys Asn Asp Gly Thr Cys Ser Asn His Pro Val Asp Phe Tyr Arg Cys Thr Cys 920 Pro Tyr Gly Phe Lys Gly Gln Asp Cys Glu Glu Pro Ile His Ala Cys 930 Ile Ser Asn Pro Cys Gln Asn Gly Gly Thr Cys His Leu Lys Asp Gly Glu Glu Asn Thr His Trp Cys Val Cys Pro Glu Gly Phe Glu Gly Asp Glu Cys Glu Ile Asn Ile Asp Asp Cys Glu Asp Asn Asp Cys Glu Asn 980 Asn Ser Thr Cys Val Asp Gly Ile Asn Asn Tyr Thr Cys Leu Cys Ser 1000 Pro Glu Tyr Thr Gly Glu Leu Cys Glu Asp Lys Leu Asp Phe Cys Ala 1010 Ser Glu Leu Asn Leu Cys Gln His Asp Ser Lys Cys Ile Leu Thr Ala

1035

1025

1030

- Lys Gly Phe Met Cys Glu Cys Thr Pro Gly Tyr Thr Gly Glu His Cys 1045 1050 1055
- Glu Val Asp Phe Asp Asp Cys Glu Asp Asn Lys Cys Lys Asn Gly Ala 1060 1065 1070
- Gln Cys Ile Asp Ala Val Asn Gly Tyr Thr Cys Val Cys Pro Glu Gly 1075 1080 1085
- Tyr Ser Gly Leu Phe Cys Glu Phe Ser Pro Pro Met Val Leu Pro Arg 1090 1095 1100
- Thr Ser Pro Cys Asp His Tyr Asp Cys Ala Asn Gly Ala Gln Cys Val 1105 1110 1115 1120
- Val Lys Asp Thr Asp Pro Val Cys Gln Cys Leu Pro Gly Tyr Glu Gly
  1125 1130 1135
- Val His Cys Glu Arg Leu Val Ser Val Asn Phe Ile Asn Arg Glu Ser 1140 1145 1150
- Phe Leu Gln Ile Pro Ser Asn Leu Ile Thr Glu Gln Ala Asn Ile Ser 1155 1160 1165
- Leu Gln Ile Ala Thr Asp Glu Asp Asn Gly Val Leu Leu Tyr Lys Gly
  1170 1180
- Asp Asn Glu His Ile Ala Val Glu Leu Tyr Arg Gly Arg Leu Arg Val 1185 1190 1195 1200
- Ser Tyr Asp Ser Gly Ser Tyr Pro Pro Ser Ala Ile Tyr Ser Val Glu 1205 1210 1215
- Thr Ile Asn Asp Gly Ser Phe His Val Val Glu Leu Val Ala Lys Asp 1220 1225 1230
- Gln Ser Leu Ser Leu Ser Ile Asp Gly Gly Ser Pro Lys Ser Ile Asn 1235 1240 1245
- Thr Ala Asn Ser Pro Ser Pro Val Pro Ser Pro Ala Pro Leu Tyr Leu 1250 1255 1260
- Gly Gly Leu Pro Gln Gln Ser Gly Leu Ala Ser Leu Arg Gln Gly Ser 1265 1270 1275 1280
- Gly Arg Asn Gly Ser Ser Phe His Gly Cys Ile Arg Asn Leu Tyr Ile 1285 1290 1295
- Asn Asp Gln Leu Gln Asp Leu Thr Gln Phe Leu Leu Gln Glu Gly Val 1300 1305 1310
- Val Pro Gly Cys Gln Pro Cys Gln Arg Ser Met Cys Ala His Gly Gln 1315 1320 1325
- Cys His Ala Thr Gly Gln Ser Ser Phe Ser Cys Glu Cys Glu Ala Gly 1330 1340

Trp Thr Gly Pro Leu Cys Asp Gln Gln Val Asn Asn Pro Cys Asp Gly 1345 1350 1355 1360

Asn Lys Cys Ile His Gly Ser Cys Met Ala Ile Asn Ser Tyr Ser Tyr 1365 1370 1375

Ser Cys Arg Cys Leu Pro Gly Phe Ala Gly Val Leu Cys Asp Glu Glu 1380 1385 1390

Glu Gln Leu Ser Pro Cys Gln Tyr Ile Ala Cys Lys Tyr Gly Arg Cys 1395 1400 1405

Arg Val Ser Gly Leu Gly Lys Ala Tyr Cys Glu Cys Asn Ser Gly Tyr 1410 1415 1420

Thr Gly Gln Ser Cys Asp Arg Glu Met Ala Cys Arg Gly Glu Arg Val 1425 1430 1435 1440

Arg Asp His Tyr Gln Thr Gln Gln Gly Tyr Ala Ala Cys Gln Ser Thr 1445 1450 1455

Glu Lys Val Ser Arg Leu Glu Cys Arg Gly Ser Cys Gly Asp Gly Thr 1460 1465 1470

Ser Cys Cys Ala Pro Leu Arg Ser Lys Arg Arg Lys Tyr Thr Phe Gln
1475 1480 1485

Cys Thr Asp Gly Ser Ser Phe Val Gln Glu Val Glu Lys Val Val Lys 1490 1495 1500

Cys Gly Cys Thr Lys Cys Pro Ser 1505 1510

<210> 33

<211> 727

<212> PRT

<213> Mus musculus

<400> 33

Met Trp Trp Arg Asn Ala Pro Ala Thr Glu Glu Asn Gly Pro Phe Leu
1 5 10 15

Arg Lys Cys Phe Ala Ile Ser Leu Leu Val Ile Cys Ile Ile Ile Ser 20 25 30

Ile Gly Ile Phe Tyr Gly Phe Val Ala Asn His Gln Val Arg Thr Arg
35 40 45

Ile Lys Arg Ser Arg Lys Leu Ala Asp Ser Asn Phe Lys Asp Leu Arg
50 55 60

Thr Leu Leu Asn Glu Thr Pro Glu Gln Ile Lys Tyr Ile Leu Ala Gln 65 70 75 80

Tyr Asn Thr Thr Lys Asp Lys Ala Phe Thr Asp Leu Asn Ser Ile Asn

85	90	95

Ser	Val	Leu	Gly 100	Gly	Gly	Ile	Leu	Asp 105	Arg	Leu	Arg	Pro	Asn 110		Ile
Pro	Val	Leu 115	Asp	Glu	Ile	Lys	Ser 120	Met	Ala	Thr	Ala	Ile 125		Glu	Thr
Lys	Glu 130	Ala	Leu	Glu	Asn	Met 135	Asn	Ser	Thr	Leu	Lys 140	Ser	Leu	His	Glr
Gln 145	Ser	Thr	Gln	Leu	Ser 150	Ser	Ser	Leu	Thr	Ser 155	Val	Lys	Thr	Ser	Leu 160
Arg	Ser	Ser	Leu	Asn 165	Asp	Pro	Leu	Cys	Leu 170	Val	His	Pro	Ser	Ser 175	Glu
Thr	Cys	Asn	Ser 180	Ile	Arg	Leu	Ser	Leu 185	Ser	Gln	Leu	Asn	Ser 190	Asn	Pro
Glu	Leu	Arg 195	Gln	Leu	Pro	Pro	Val 200	Asp	Ala	Glu	Leu	Asp 205	Asn	Val	Asn
	210					215	Asp				220			_	
225					230		Arg			235					240
				245			Leu		250					255	Ī
			260				Ile	265					270		
		275					Ser 280					285			
	290					295	Trp				300				
305					310		Ile			315					320
				325			His		330					335	·
			340				Phe	345					350		
		355					Met 360					365			
	370					375	Leu				380				
GIU	Leu	rne	Arg	۷al	ьeu	Asp	Thr	Pro	Tyr	Leu	Leu	Asn	Glu	Asp	Trp

Glu Tyr Tyr Leu Ser Gly Lys Leu Phe Asn Lys Ser Lys Met Lys Leu 405 410 415

Thr Phe Glu Gln Val Tyr Ser Asp Cys Lys Lys Asn Arg Gly Thr Tyr 420 425 430

Gly Thr Leu His Leu Gln Asn Ser Phe Asn Ile Ser Glu His Leu Asn 435 440 445

Ile Asn Glu His Thr Gly Ser Ile Ser Ser Glu Leu Glu Ser Leu Lys 450 455 460

Val Asn Leu Asn Ile Phe Leu Leu Gly Ala Ala Gly Arg Lys Asn Leu 465 470 475 480

Gln Asp Phe Ala Ala Cys Gly Ile Asp Arg Met Asn Tyr Asp Ser Tyr 485 490 495

Leu Ala Gln Thr Gly Lys Ser Pro Ala Gly Val Asn Leu Leu Ser Phe 500 505 510

Ala Tyr Asp Leu Glu Ala Lys Ala Asn Ser Leu Pro Pro Gly Asn Leu 515 520 525

Arg Asn Ser Leu Lys Arg Asp Ala Gln Thr Ile Lys Thr Ile His Gln 530 535 540

Gln Arg Val Leu Pro Ile Glu Gln Ser Leu Ser Thr Leu Tyr Gln Ser 545 550 555 560

Val Lys Ile Leu Gln Arg Thr Gly Asn Gly Leu Leu Glu Arg Val Thr 565 570 575

Arg Ile Leu Ala Ser Leu Asp Phe Ala Gln Asn Phe Ile Thr Asn Asn 580 585 590

Thr Ser Ser Val Ile Ile Glu Glu Thr Lys Lys Tyr Gly Arg Thr Ile 595 600 605

Ile Gly Tyr Phe Glu His Tyr Leu Gln Trp Ile Glu Phe Ser Ile Ser 610 620

Glu Lys Val Ala Ser Cys Lys Pro Val Ala Thr Ala Leu Asp Thr Ala 625 630 635 640

Val Asp Val Phe Leu Cys Ser Tyr Ile Ile Asp Pro Leu Asn Leu Phe 645 650 655

Trp Phe Gly Ile Gly Lys Ala Thr Val Phe Leu Leu Pro Ala Leu Ile 660 665 670

Phe Ala Val Lys Leu Ala Lys Tyr Tyr Arg Arg Met Asp Ser Glu Asp 675 680 685

Val Tyr Asp Asp Val Glu Thr Ile Pro Met Lys Asn Met Glu Asn Gly

690 695 700

Asn Asn Gly Tyr His Lys Asp His Val Tyr Gly Ile His Asn Pro Val 705 710 715 720

Met Thr Ser Pro Ser Gln His
725

<210> 34

<211> 856

<212> PRT

<213> Homo sapiens

<400> 34

Met Ala Leu Val Leu Gly Ser Leu Leu Leu Leu Gly Leu Cys Gly Asn 1 5 10 15

Ser Phe Ser Gly Gly Gln Pro Ser Ser Thr Asp Ala Pro Lys Ala Trp
20 25 30

Asn Tyr Glu Leu Pro Ala Thr Asn Tyr Glu Thr Gln Asp Ser His Lys
35 40 45

Ala Gly Pro Ile Gly Ile Leu Phe Glu Leu Val His Ile Phe Leu Tyr
50 55 60

Val Val Gln Pro Arg Asp Phe Pro Glu Asp Thr Leu Arg Lys Phe Leu 65 70 75 80

Gln Lys Ala Tyr Glu Ser Lys Ile Asp Tyr Asp Lys Ile Val Tyr Tyr 85 90 95

Glu Ala Gly Ile Ile Leu Cys Cys Val Leu Gly Leu Leu Phe Ile Ile 100 105 110

Leu Met Pro Leu Val Gly Tyr Phe Phe Cys Met Cys Arg Cys Cys Asn 115 120 125

Lys Cys Gly Glu Met His Gln Arg Gln Lys Glu Asn Gly Pro Phe 130 135 140

Leu Arg Lys Cys Phe Ala Ile Ser Leu Leu Val Ile Cys Ile Ile Ile 145 150 155 160

Ser Ile Gly Ile Phe Tyr Gly Phe Val Ala Asn His Gln Val Arg Thr 165 170 175

Arg Ile Lys Arg Ser Arg Lys Leu Ala Asp Ser Asn Phe Lys Asp Leu 180 185 190

Arg Thr Leu Leu Asn Glu Thr Pro Glu Gln Ile Lys Tyr Ile Leu Ala 195 200 205

Gln Tyr Asn Thr Thr Lys Asp Lys Ala Phe Thr Asp Leu Asn Ser Ile 210 215 220

Asn Ser Val Leu Gly Gly Gly Ile Leu Asp Arg Leu Arg Pro Asn Ile 225 230 Ile Pro Val Leu Asp Glu Ile Lys Ser Met Ala Thr Ala Ile Lys Glu 245 250 Thr Lys Glu Ala Leu Glu Asn Met Asn Ser Thr Leu Lys Ser Leu His 265 Gln Gln Ser Thr Gln Leu Ser Ser Leu Thr Ser Val Lys Thr Ser 280 Leu Arg Ser Ser Leu Asn Asp Pro Leu Cys Leu Val His Pro Ser Ser Glu Thr Cys Asn Ser Ile Arg Leu Ser Leu Ser Gln Leu Asn Ser Asn 310 315 Pro Glu Leu Arg Gln Leu Pro Pro Val Asp Ala Glu Leu Asp Asn Val 325 Asn Asn Val Leu Arg Thr Asp Leu Asp Gly Leu Val Gln Gln Gly Tyr 345 Gln Ser Leu Asn Asp Ile Pro Asp Arg Val Gln Arg Gln Thr Thr Val Val Ala Gly Ile Lys Arg Val Leu Asn Ser Ile Gly Ser Asp Ile Asp Asn Val Thr Gln Arg Leu Pro Ile Gln Asp Ile Leu Ser Ala Phe 390 395 Ser Val Tyr Val Asn Asn Thr Glu Ser Tyr Ile His Arg Asn Leu Pro 405 410 Thr Leu Glu Glu Tyr Asp Ser Tyr Trp Trp Leu Gly Gly Leu Val Ile 425 Cys Ser Leu Leu Thr Leu Ile Val Ile Phe Tyr Tyr Leu Gly Leu Leu 435 Cys Gly Val Cys Gly Tyr Asp Arg His Ala Thr Pro Thr Thr Arg Gly Cys Val Ser Asn Thr Gly Gly Val Phe Leu Met Val Gly Val Gly Leu 470 475 Ser Phe Leu Phe Cys Trp Ile Leu Met Ile Ile Val Val Leu Thr Phe 490 Val Phe Gly Ala Asn Val Glu Lys Leu Ile Cys Glu Pro Tyr Thr Ser 505 Lys Glu Leu Phe Arg Val Leu Asp Thr Pro Tyr Leu Leu Asn Glu Asp 515 520

Trp Glu Tyr Tyr Leu Ser Gly Lys Leu Phe Asn Lys Ser Lys Met Lys 535 Leu Thr Phe Glu Gln Val Tyr Ser Asp Cys Lys Lys Asn Arg Gly Thr 550 555 Tyr Gly Thr Leu His Leu Gln Asn Ser Phe Asn Ile Ser Glu His Leu 570 Asn Ile Asn Glu His Thr Gly Ser Ile Ser Ser Glu Leu Glu Ser Leu Lys Val Asn Leu Asn Ile Phe Leu Leu Gly Ala Ala Gly Arg Lys Asn 600 Leu Gln Asp Phe Ala Ala Cys Gly Ile Asp Arg Met Asn Tyr Asp Ser 615 Tyr Leu Ala Gln Thr Gly Lys Ser Pro Ala Gly Val Asn Leu Leu Ser 630 635 Phe Ala Tyr Asp Leu Glu Ala Lys Ala Asn Ser Leu Pro Pro Gly Asn 645 650 Leu Arg Asn Ser Leu Lys Arg Asp Ala Gln Thr Ile Lys Thr Ile His Gln Gln Arg Val Leu Pro Ile Glu Gln Ser Leu Ser Thr Leu Tyr Gln 680 Ser Val Lys Ile Leu Gln Arg Thr Gly Asn Gly Leu Leu Glu Arg Val 695 Thr Arg Ile Leu Ala Ser Leu Asp Phe Ala Gln Asn Phe Ile Thr Asn 710 715 Asn Thr Ser Ser Val Ile Ile Glu Glu Thr Lys Lys Tyr Gly Arg Thr Ile Ile Gly Tyr Phe Glu His Tyr Leu Gln Trp Ile Glu Phe Ser Ile 740 Ser Glu Lys Val Ala Ser Cys Lys Pro Val Ala Thr Ala Leu Asp Thr Ala Val Asp Val Phe Leu Cys Ser Tyr Ile Ile Asp Pro Leu Asn Leu 770 780 Phe Trp Phe Gly Ile Gly Lys Ala Thr Val Phe Leu Leu Pro Ala Leu Ile Phe Ala Val Lys Leu Ala Lys Tyr Tyr Arg Arg Met Asp Ser Glu 810 Asp Val Tyr Asp Asp Val Glu Thr Ile Pro Met Lys Asn Met Glu Asn 820 825

Gly Asn Asn Gly Tyr His Lys Asp His Val Tyr Gly Ile His Asn Pro 835 840 845

Val Met Thr Ser Pro Ser Gln His 850 855

<210> 35

<211> 865

<212> PRT

<213> Homo sapiens

<400> 35

Met Ala Leu Val Leu Gly Ser Leu Leu Leu Leu Gly Leu Cys Gly Asn
1 5 10 15

Ser Phe Ser Gly Gly Gln Pro Ser Ser Thr Asp Ala Pro Lys Ala Trp
20 25 30

Asn Tyr Glu Leu Pro Ala Thr Asn Tyr Glu Thr Gln Asp Ser His Lys
35 40 45

Ala Gly Pro Ile Gly Ile Leu Phe Glu Leu Val His Ile Phe Leu Tyr
50 55 60

Val Val Gln Pro Arg Asp Phe Pro Glu Asp Thr Leu Arg Lys Phe Leu
65 70 75 80

Gln Lys Ala Tyr Glu Ser Lys Ile Asp Tyr Asp Lys Pro Glu Thr Val 85 90 95

Ile Leu Gly Leu Lys Ile Val Tyr Tyr Glu Ala Gly Ile Ile Leu Cys
100 105 110

Cys Val Leu Gly Leu Leu Phe Ile Ile Leu Met Pro Leu Val Gly Tyr 115 120 125

Phe Phe Cys Met Cys Arg Cys Cys Asn Lys Cys Gly Glu Met His 130 135 140

Gln Arg Gln Lys Glu Asn Gly Pro Phe Leu Arg Lys Cys Phe Ala Ile 145 150 155 160

Ser Leu Leu Val Ile Cys Ile Ile Ile Ser Ile Gly Ile Phe Tyr Gly 165 170 175

Phe Val Ala Asn His Gln Val Arg Thr Arg Ile Lys Arg Ser Arg Lys
180 185 190

Leu Ala Asp Ser Asn Phe Lys Asp Leu Arg Thr Leu Leu Asn Glu Thr
195 200 205

Pro Glu Gln Ile Lys Tyr Ile Leu Ala Gln Tyr Asn Thr Thr Lys Asp 210 215 220

Lys Ala Phe Thr Asp Leu Asn Ser Ile Asn Ser Val Leu Gly Gly 225 230 235 240

Ile Leu Asp Arg Leu Arg Pro Asn Ile Ile Pro Val Leu Asp Glu Ile 250 Lys Ser Met Ala Thr Ala Ile Lys Glu Thr Lys Glu Ala Leu Glu Asn Met Asn Ser Thr Leu Lys Ser Leu His Gln Gln Ser Thr Gln Leu Ser 280 Ser Ser Leu Thr Ser Val Lys Thr Ser Leu Arg Ser Ser Leu Asn Asp 295 300 Pro Leu Cys Leu Val His Pro Ser Ser Glu Thr Cys Asn Ser Ile Arq 310 315 Leu Ser Leu Ser Gln Leu Asn Ser Asn Pro Glu Leu Arg Gln Leu Pro 330 Pro Val Asp Ala Glu Leu Asp Asn Val Asn Asn Val Leu Arg Thr Asp 345 Leu Asp Gly Leu Val Gln Gln Gly Tyr Gln Ser Leu Asn Asp Ile Pro 360 Asp Arg Val Gln Arg Gln Thr Thr Thr Val Val Ala Gly Ile Lys Arg 375 Val Leu Asn Ser Ile Gly Ser Asp Ile Asp Asn Val Thr Gln Arg Leu 390 395 Pro Ile Gln Asp Ile Leu Ser Ala Phe Ser Val Tyr Val Asn Asn Thr 405 Glu Ser Tyr Ile His Arg Asn Leu Pro Thr Leu Glu Glu Tyr Asp Ser 420 425 Tyr Trp Trp Leu Gly Gly Leu Val Ile Cys Ser Leu Leu Thr Leu Ile 435 Val Ile Phe Tyr Tyr Leu Gly Leu Leu Cys Gly Val Cys Gly Tyr Asp Arg His Ala Thr Pro Thr Thr Arg Gly Cys Val Ser Asn Thr Gly Gly 465 470 475 Val Phe Leu Met Val Gly Val Gly Leu Ser Phe Leu Phe Cys Trp Ile 485 Leu Met Ile Ile Val Val Leu Thr Phe Val Phe Gly Ala Asn Val Glu 505 Lys Leu Ile Cys Glu Pro Tyr Thr Ser Lys Glu Leu Phe Arg Val Leu 515 Asp Thr Pro Tyr Leu Leu Asn Glu Asp Trp Glu Tyr Tyr Leu Ser Gly 535 540

Lys Leu Phe Asn Lys Ser Lys Met Lys Leu Thr Phe Glu Gln Val Tyr 550 555 Ser Asp Cys Lys Lys Asn Arg Gly Thr Tyr Gly Thr Leu His Leu Gln 570 Asn Ser Phe Asn Ile Ser Glu His Leu Asn Ile Asn Glu His Thr Gly 580 585 Ser Ile Ser Ser Glu Leu Glu Ser Leu Lys Val Asn Leu Asn Ile Phe 600 Leu Leu Gly Ala Ala Gly Arg Lys Asn Leu Gln Asp Phe Ala Ala Cys 610 615 Gly Ile Asp Arg Met Asn Tyr Asp Ser Tyr Leu Ala Gln Thr Gly Lys 630 635 Ser Pro Ala Gly Val Asn Leu Leu Ser Phe Ala Tyr Asp Leu Glu Ala 650 Lys Ala Asn Ser Leu Pro Pro Gly Asn Leu Arg Asn Ser Leu Lys Arg 660 Asp Ala Gln Thr Ile Lys Thr Ile His Gln Gln Arg Val Leu Pro Ile 680 Glu Gln Ser Leu Ser Thr Leu Tyr Gln Ser Val Lys Ile Leu Gln Arg 690 695 Thr Gly Asn Gly Leu Leu Glu Arg Val Thr Arg Ile Leu Ala Ser Leu 705 Asp Phe Ala Gln Asn Phe Ile Thr Asn Asn Thr Ser Ser Val Ile Ile 730 Glu Glu Thr Lys Lys Tyr Gly Arg Thr Ile Ile Gly Tyr Phe Glu His Tyr Leu Gln Trp Ile Glu Phe Ser Ile Ser Glu Lys Val Ala Ser Cys Lys Pro Val Ala Thr Ala Leu Asp Thr Ala Val Asp Val Phe Leu Cys 770 775 Ser Tyr Ile Ile Asp Pro Leu Asn Leu Phe Trp Phe Gly Ile Gly Lys 785 Ala Thr Val Phe Leu Leu Pro Ala Leu Ile Phe Ala Val Lys Leu Ala 805 810 Lys Tyr Tyr Arg Arg Met Asp Ser Glu Asp Val Tyr Asp Asp Val Glu 820 825 Thr Ile Pro Met Lys Asn Met Glu Asn Gly Asn Asn Gly Tyr His Lys 835 840 845

Asp His Val Tyr Gly Ile His Asn Pro Val Met Thr Ser Pro Ser Gln 855 860 His 865 <210> 36 <211> 857 <212> PRT <213> Rattus norvegicus <400> 36 Met Ala Leu Val Phe Ser Val Leu Leu Leu Gly Leu Cys Gly Lys Met Ala Ser Gly Gly Gln Pro Ala Phe Asp Asn Thr Pro Gly Ala Leu Asn Tyr Glu Leu Pro Thr Thr Glu Tyr Glu Thr Gln Asp Thr Phe Asn Ala Gly Ile Ile Asp Pro Leu Tyr Gln Met Val His Ile Phe Leu Asn Val Val Gln Pro Asn Asp Phe Pro Gln Asp Leu Val Lys Lys Leu Ile 65 70 Gln Lys Arg Phe Asp Ile Ser Val Asp Thr Lys Glu Val Ala Ile Tyr Glu Ile Gly Val Leu Ile Cys Val Ile Leu Gly Leu Leu Phe Ile Phe 105 Leu Met Pro Leu Val Gly Phe Phe Phe Cys Met Cys Arg Cys Cys Asn 115 Lys Cys Gly Glu Met His Gln Arg Gln Lys Gln Asn Glu Ser Cys 135 Arg Arg Lys Cys Leu Ala Ile Ser Leu Leu Ile Cys Leu Leu Met 150 155 Ser Leu Gly Ile Ala Phe Gly Phe Val Ala Asn Gln Gln Thr Arg Thr Arg Ile Gln Arg Thr Gln Lys Leu Ala Glu Ser Asn Tyr Arg Asp Leu 185 Arg Ala Leu Leu Thr Glu Ala Pro Lys Gln Ile Asp Tyr Ile Leu Gly 195 Gln Tyr Asn Thr Thr Lys Asn Lys Ala Phe Ser Asp Leu Asp Ser Ile

Asp Ser Val Leu Gly Gly Arg Ile Lys Gly Gln Leu Lys Pro Lys Val

225 230	235	240
---------	-----	-----

Thr Pro Val Leu Glu Glu Ile Lys Ala Met Ala Thr Ala Ile Arg Gln 245 250 255

Thr Lys Asp Ala Leu Gln Asn Met Ser Ser Ser Leu Lys Ser Leu Arg 260 265 270

Asp Ala Ser Thr Gln Leu Ser Thr Asn Leu Thr Ser Val Arg Asn Ser 275 280 285

Ile Glu Asn Ser Leu Asn Ser Asn Asp Cys Ala Ser Asp Pro Ala Ser 290 295 300

Lys Ile Cys Asp Ser Leu Arg Pro Gln Leu Ser Asn Leu Gly Ser Asn 305 310 315 320

His Asn Gly Ser Gln Leu Pro Ser Val Asp Arg Glu Leu Asn Thr Val
325 330 335

Asn Asp Val Asp Arg Thr Asp Leu Glu Ser Leu Val Lys Arg Gly Tyr 340 345 350

Met Ser Ile Asp Glu Ile Pro Asn Met Ile Gln Asn Gln Thr Gly Asp 355 360 365

Val Ile Lys Asp Val Lys Lys Thr Leu Asp Ser Val Ser Ser Lys Val 370 375 380

Lys Asn Met Ser Gln Ser Ile Pro Val Glu Glu Val Leu Leu Gln Phe 385 390 395 400

Ser His Tyr Leu Asn Asp Ser Asn Arg Tyr Ile His Glu Ser Leu Pro 405 410 415

Arg Val Glu Glu Tyr Asp Ser Tyr Trp Trp Leu Gly Gly Leu Ile Val 420 425 430

Cys Phe Leu Leu Thr Leu Ile Val Thr Phe Phe Tyr Leu Gly Leu Leu 435 440 445

Cys Gly Val Phe Gly Tyr Asp Lys Arg Ala Thr Pro Thr Arg Arg Gly
450 455 460

Cys Val Ser Asn Thr Gly Gly Ile Phe Leu Met Ala Gly Val Gly Phe
465 470 475 480

Ser Phe Leu Phe Cys Trp Ile Leu Met Ile Leu Val Val Leu Thr Phe 485 490 495

Val Val Gly Ala Asn Val Glu Lys Leu Leu Cys Glu Pro Tyr Glu Asn 500 505 510

Lys Lys Leu Leu Gln Val Leu Asp Thr Pro Tyr Leu Leu Asn Asp Gln 515 520 525

Trp Gln Phe Tyr Leu Ser Gly Ile Leu Leu Lys Asn Pro Asp Ile Asn

Met 545	Thr	Phe	Glu	Gln	Val 550	Tyr	Arg	Asp	Cys	Lys 555	Arg	Gly	Arg	Gly	Va. 560
Tyr	Ala	Thr	Phe	Gln 565	Leu	Glu	Asn	Val	Phe 570	Asn	Ile	Thr	Glu	Asn 575	
Asn	Ile	Glu	Arg 580	Leu	Ser	Glu	Asp	Ile 585	Val	Lys	Glu	Leu	Glu 590	Lys	Lei
Asn	Val	Asn 595	Ile	Asp	Ser	Ile	Glu 600	Leu	Leu	Asp	Lys	Thr 605	Gly	Arg	Lys
Ser	Leu 610	Glu	Asp	Phe	Ala	Gln 615	Ser	Gly	Ile	Asp	Arg 620	Ile	Asn	Tyr	Sei
Met 625	Tyr	Leu	Gln	Glu	Ala 630	Glu	Lys	Pro	Pro	Thr 635	Lys	Val	Asp	Leu	Leu 640
Thr	Phe	Ala	Ser	Phe 645	Leu	Glu	Thr	Glu	Ala 650	Asn	Gln	Leu	Pro	Asp 655	Gly
Asn	Leu	Lys	Gln 660	Ala	Phe	Leu	Met	Asp 665	Ala	Gln	Asn	Ile	Arg 670	Ala	Ile
His	Gln	Gln 675	His	Val	Pro	Pro	Val 680	Gln	Gln	Ser	Leu	Asn 685	Ser	Leu	Lys
Gln	Ser 690	Val	Trp	Ala	Leu	Lys 695	Gln	Thr	Ser	Ser	Lys 700	Leu	Pro	Glu	Glu
Val 705	Lys	Lys	Val	Leu	Ala 710	Ser	Leu	Asp	Ser	Ala 715	Gln	His	Phe	Leu	Thr 720
Ser	Asn	Leu	Ser	Ser 725	Ile	Val	Ile	Gly	Glu 730	Thr	Lys	Lys	Phe	Gly 735	Arg
Thr	Ile	Ile	Gly 740	Tyr	Phe	Glu	His	Tyr 745	Leu	Gln	Trp	Val	Leu 750	Tyr	Ala
Ile	Thr	Glu 755	Lys	Met	Thr	Ser	Cys 760	Lys	Pro	Met	Ile	Thr 765	Ala	Met	Asp
Ser	Ala 770	Val	Asn	Gly	Ile	Leu 775	Cys	Ser	Tyr	Val	Ala 780	Asp	Pro	Leu	Asn
Leu 785	Phe	Trp	Phe	Gly	Ile 790	Gly	Lys	Ala	Thr	Met 795	Leu	Leu	Leu	Pro	Ala 800
Val	Ile	Ile	Ala	Ile 805	Lys	Leu	Ala	Lys	Tyr 810	Tyr	Arg	Arg	Met	Asp 815	Ser
Glu	Asp	Val	Tyr 820	Asp	Asp	Val	Glu	Thr 825	Val	Pro	Met	Lys	Asn 830	Leu	Glu

Asn Gly Ser Asn Gly Tyr His Lys Asp His Leu Tyr Gly Val His Asn

Pro Val Met Thr Ser Pro Ser Arg Tyr 850 855

<210> 37

<211> 867

<212> PRT

<213> Mus musculus

<400> 37

Met Ala Leu Val Phe Ser Ala Leu Leu Leu Leu Gly Leu Cys Gly Lys

1 10 15

Ile Ser Ser Glu Gly Gln Pro Ala Phe His Asn Thr Pro Gly Ala Met 20 25 30

Asn Tyr Glu Leu Pro Thr Thr Lys Tyr Glu Thr Gln Asp Thr Phe Asn 35 40 45

Ala Gly Ile Val Gly Pro Leu Tyr Lys Met Val His Ile Phe Leu Ser 50 55 60

Val Val Gln Pro Asn Asp Phe Pro Leu Asp Leu Ile Lys Lys Leu Ile
65 70 75 80

Gln Asn Lys Lys Phe Asp Ile Ser Val Asp Ser Lys Glu Pro Glu Ile 85 90 95

Ile Val Leu Ala Leu Lys Ile Ala Leu Tyr Glu Ile Gly Val Leu Ile 100 105 110

Cys Ala Ile Leu Gly Leu Leu Phe Ile Ile Leu Met Pro Leu Val Gly 115 120 125

Cys Phe Phe Cys Met Cys Arg Cys Cys Asn Lys Cys Gly Glu Met 130 135 140

His Gln Arg Gln Lys Gln Asn Ala Pro Cys Arg Arg Lys Cys Leu Gly
145 150 155 160

Leu Ser Leu Leu Val Ile Cys Leu Leu Met Ser Leu Gly Ile Ile Tyr 165 170 175

Gly Phe Val Ala Asn Gln Gln Thr Arg Thr Arg Ile Lys Gly Thr Gln
180 185 190

Lys Leu Ala Lys Ser Asn Phe Arg Asp Phe Gln Thr Leu Leu Thr Glu 195 200 205

Thr Pro Lys Gln Ile Asp Tyr Val Val Glu Gln Tyr Thr Asn Thr Lys 210 215 220

Asn Lys Ala Phe Ser Asp Leu Asp Gly Ile Gly Ser Val Leu Gly Gly 225 230 235 240

Arg Ile Lys Asp Gln Leu Lys Pro Lys Val Thr Pro Val Leu Glu Glu 245 250 Ile Lys Ala Met Ala Thr Ala Ile Lys Gln Thr Lys Asp Ala Leu Gln 260 265 Asn Met Ser Ser Leu Lys Ser Leu Gln Asp Ala Ala Thr Gln Leu 280 Asn Thr Asn Leu Ser Ser Val Arg Asn Ser Ile Glu Asn Ser Leu Ser 295 Ser Ser Asp Cys Thr Ser Asp Pro Ala Ser Lys Ile Cys Asp Ser Ile Arg Pro Ser Leu Ser Ser Leu Gly Ser Ser Leu Asn Ser Ser Gln Leu 325 330 Pro Ser Val Asp Arg Glu Leu Asn Thr Val Thr Glu Val Asp Lys Thr 340 345 Asp Leu Glu Ser Leu Val Lys Arg Gly Tyr Thr Thr Ile Asp Glu Ile 360 Pro Asn Thr Ile Gln Asn Gln Thr Val Asp Val Ile Lys Asp Val Lys 370 Asn Thr Leu Asp Ser Ile Ser Ser Asn Ile Lys Asp Met Ser Gln Ser 390 Ile Pro Ile Glu Asp Met Leu Leu Gln Val Ser His Tyr Leu Asn Asn 405 410 Ser Asn Arg Tyr Leu Asn Gln Glu Leu Pro Lys Leu Glu Glu Tyr Asp 420 Ser Tyr Trp Trp Leu Gly Gly Leu Ile Val Cys Phe Leu Leu Thr Leu Ile Val Thr Phe Phe Phe Leu Gly Leu Leu Cys Gly Val Phe Gly Tyr 450 Asp Lys His Ala Thr Pro Thr Arg Arg Gly Cys Val Ser Asn Thr Gly 470 Gly Ile Phe Leu Met Ala Gly Val Gly Phe Gly Phe Leu Phe Cys Trp 490 Ile Leu Met Ile Leu Val Val Leu Thr Phe Val Val Gly Ala Asn Val 500 505 Glu Lys Leu Cys Glu Pro Tyr Glu Asn Lys Lys Leu Leu Gln Val Leu Asp Thr Pro Tyr Leu Leu Lys Glu Gln Trp Gln Phe Tyr Leu Ser 530

Gly Met Leu Phe Asn Asn Pro Asp Ile Asn Met Thr Phe Glu Gln Val 545 Tyr Arg Asp Cys Lys Arg Gly Arg Gly Ile Tyr Ala Ala Phe Gln Leu 565 570 Glu Asn Val Val Asn Val Ser Asp His Phe Asn Ile Asp Gln Ile Ser 585 Glu Asn Ile Asn Thr Glu Leu Glu Asn Leu Asn Val Asn Ile Asp Ser Ile Glu Leu Leu Asp Asn Thr Gly Arg Lys Ser Leu Glu Asp Phe Ala His Ser Gly Ile Asp Thr Ile Asp Tyr Ser Thr Tyr Leu Lys Glu Thr 630 Glu Lys Ser Pro Thr Glu Val Asn Leu Leu Thr Phe Ala Ser Thr Leu 645 650 Glu Ala Lys Ala Asn Gln Leu Pro Glu Gly Lys Pro Lys Gln Ala Phe 665 Leu Leu Asp Val Gln Asn Ile Arg Ala Ile His Gln His Leu Leu Pro 675 Pro Val Gln Gln Ser Leu Asn Thr Leu Arg Gln Ser Val Trp Thr Leu 695 Gln Gln Thr Ser Asn Lys Leu Pro Glu Lys Val Lys Lys Ile Leu Ala 710 715 Ser Leu Asp Ser Val Gln His Phe Leu Thr Asn Asn Val Ser Leu Ile 730 Val Ile Gly Glu Thr Lys Lys Phe Gly Lys Thr Ile Leu Gly Tyr Phe Glu His Tyr Leu His Trp Val Phe Tyr Ala Ile Thr Glu Lys Met Thr Ser Cys Lys Pro Met Ala Thr Ala Met Asp Ser Ala Val Asn Gly Ile Leu Cys Gly Tyr Val Ala Asp Pro Leu Asn Leu Phe Trp Phe Gly Ile 790 795 Gly Lys Ala Thr Val Leu Leu Pro Ala Val Ile Ile Ala Ile Lys Leu Ala Lys Tyr Tyr Arg Arg Met Asp Ser Glu Asp Val Tyr Asp Asp Val Glu Thr Val Pro Met Lys Asn Leu Glu Ile Gly Ser Asn Gly Tyr 835 840

His Lys Asp His Leu Tyr Gly Val His Asn Pro Val Met Thr Ser Pro 850 855 860

Ser Arg Tyr 865

<210> 38

<211> 331

<212> PRT

<213> Homo sapiens

<400> 38

Met Glu Asn Pro Ser Pro Ala Ala Ala Leu Gly Lys Ala Leu Cys Ala 1 5 10 15

Leu Leu Leu Ala Thr Leu Gly Ala Ala Gly Gln Pro Leu Gly Gly Glu 20 25 30

Ser Ile Cys Ser Ala Arg Ala Pro Ala Lys Tyr Ser Ile Thr Phe Thr 35 40 45

Gly Lys Trp Ser Gln Thr Ala Phe Pro Lys Gln Tyr Pro Leu Phe Arg
50 55 60

Pro Pro Ala Gln Trp Ser Ser Leu Leu Gly Ala Ala His Ser Ser Asp
65 70 75 80

Tyr Ser Met Trp Arg Lys Asn Gln Tyr Val Ser Asn Gly Leu Arg Asp 85 90 95

Phe Ala Glu Arg Gly Glu Ala Trp Ala Leu Met Lys Glu Ile Glu Ala 100 105 110

Ala Gly Glu Ala Leu Gln Ser Val His Ala Val Phe Ser Ala Pro Ala 115 120 125

Val Pro Ser Gly Thr Gly Gln Thr Ser Ala Glu Leu Glu Val Gln Arg 130 135 140

Arg His Ser Leu Val Ser Phe Val Val Arg Ile Val Pro Ser Pro Asp 145 150 155 160

Trp Phe Val Gly Val Asp Ser Leu Asp Leu Cys Asp Gly Asp Arg Trp
165 170 175

Arg Glu Gln Ala Ala Leu Asp Leu Tyr Pro Tyr Asp Ala Gly Thr Asp 180 185 190

Ser Gly Phe Thr Phe Ser Ser Pro Asn Phe Ala Thr Ile Pro Gln Asp 195 200 205

Thr Val Thr Glu Ile Thr Ser Ser Ser Pro Ser His Pro Ala Asn Ser 210 215 220

Phe Tyr Tyr Pro Arg Leu Lys Ala Leu Pro Pro Ile Ala Arg Val Thr 225 230 235 240

- Leu Val Arg Leu Arg Gln Ser Pro Arg Ala Phe Ile Pro Pro Ala Pro 245 250 255
- Val Leu Pro Ser Arg Asp Asn Glu Ile Val Asp Ser Ala Ser Val Pro 260 265 270
- Glu Thr Pro Leu Asp Cys Glu Val Ser Leu Trp Ser Ser Trp Gly Leu 275 280 285
- Cys Gly Gly His Cys Gly Arg Leu Gly Thr Lys Ser Arg Thr Arg Tyr
  290 295 300
- Val Arg Val Gln Pro Ala Asn Asn Gly Ser Pro Cys Pro Glu Leu Glu 305 310 315 320
- Glu Glu Ala Glu Cys Val Pro Asp Asn Cys Val 325 330
- <210> 39
- <211> 331
- <212> PRT
- <213> Homo sapiens
- <400> 39
- Met Glu Asn Pro Ser Pro Ala Ala Ala Leu Gly Lys Ala Leu Cys Ala 1 5 10 15
- Leu Leu Leu Ala Thr Leu Gly Ala Ala Gly Gln Pro Leu Gly Glu 20 25 30
- Ser Ile Cys Ser Ala Arg Ala Leu Ala Lys Tyr Ser Ile Thr Phe Thr 35 40 45
- Gly Lys Trp Ser Gln Thr Ala Phe Pro Lys Gln Tyr Pro Leu Phe Arg
  50 55 60
- Pro Pro Ala Gln Trp Ser Ser Leu Leu Gly Ala Ala His Ser Ser Asp
  65 70 75 80
- Tyr Ser Met Trp Arg Lys Asn Gln Tyr Val Ser Asn Gly Leu Arg Asp 85 90 95
- Phe Ala Glu Arg Gly Glu Ala Trp Ala Leu Met Lys Glu Ile Glu Ala 100 105 110
- Ala Gly Glu Ala Leu Gln Ser Val His Ala Val Phe Ser Ala Pro Ala 115 120 125
- Val Pro Ser Gly Thr Gly Gln Thr Ser Ala Glu Leu Glu Val Gln Arg 130 135 140
- Arg His Ser Leu Val Ser Phe Val Val Arg Ile Val Pro Ser Pro Asp 145 150 155 160
- Trp Phe Val Gly Val Asp Ser Leu Asp Leu Cys Asp Gly Asp Arg Trp

Arg Glu Gln Ala Ala Leu Asp Leu Tyr Pro Tyr Asp Ala Gly Thr Asp 180 185 190

Ser Gly Phe Thr Phe Ser Ser Pro Asn Phe Ala Thr Ile Pro Gln Asp 195 200 205

Thr Val Thr Glu Ile Thr Ser Ser Ser Pro Ser His Pro Ala Asn Ser 210 215 220

Phe Tyr Tyr Pro Arg Leu Lys Ala Leu Pro Pro Ile Ala Arg Val Thr 225 230 235 240

Leu Val Arg Leu Arg Gln Ser Pro Arg Ala Phe Ile Pro Pro Ala Pro 245 250 255

Val Leu Pro Ser Arg Asp Asn Glu Ile Val Asp Ser Ala Ser Val Pro 260 265 270

Glu Thr Pro Leu Asp Cys Glu Val Ser Leu Trp Ser Ser Trp Gly Leu 275 280 285

Cys Gly Gly His Cys Gly Arg Leu Gly Thr Lys Ser Arg Thr Arg Tyr 290 295 300

Val Arg Val Gln Pro Ala Asn Asn Gly Ser Pro Cys Pro Glu Leu Glu 305 310 315 320

Glu Glu Ala Glu Cys Val Pro Asp Asn Cys Val 325 330

<210> 40

<211> 331

<212> PRT

<213> Homo sapiens

<400> 40

Met Glu Asn Pro Ser Pro Ala Ala Ala Leu Gly Lys Ala Leu Cys Ala 1 5 10 15

Leu Leu Leu Ala Thr Leu Gly Ala Ala Gly Gln Pro Leu Gly Gly Glu
20 25 30

Ser Ile Cys Ser Ala Arg Ala Pro Ala Lys Tyr Ser Ile Thr Phe Thr 35 40 45

Gly Lys Trp Ser Gln Thr Ala Phe Pro Lys Gln Tyr Pro Leu Phe Arg
50 55 60

Pro Pro Ala Gln Trp Ser Ser Leu Leu Gly Ala Ala His Ser Ser Asp 65 70 75 80

Tyr Ser Met Trp Arg Lys Asn Gln Tyr Val Ser Asn Gly Leu Arg Asp 85 90 95

Phe Ala Glu Arg Gly Glu Ala Trp Ala Leu Met Lys Glu Ile Glu Ala 100 105 Ala Gly Glu Ala Leu Gln Ser Val His Glu Val Phe Ser Ala Pro Ala 120 Val Pro Ser Gly Thr Gly Gln Thr Ser Ala Glu Leu Glu Val Gln Arg 135 Arg His Ser Leu Val Ser Phe Val Val Arg Ile Val Pro Ser Pro Asp 160 Trp Phe Val Gly Val Asp Ser Leu Asp Leu Cys Asp Gly Asp Arg Trp Arg Glu Gln Ala Ala Leu Asp Leu Tyr Pro Tyr Asp Ala Gly Thr Asp Ser Gly Phe Thr Phe Ser Ser Pro Asn Phe Ala Thr Ile Pro Gln Asp 195 200 Thr Val Thr Glu Ile Thr Ser Ser Pro Ser His Pro Ala Asn Ser 215 Phe Tyr Tyr Pro Arg Leu Lys Ala Leu Pro Pro Ile Ala Arg Val Thr 240 Leu Leu Arg Leu Arg Gln Ser Pro Arg Ala Phe Ile Pro Pro Ala Pro 245 250 Val Leu Pro Ser Arg Asp Asn Glu Ile Val Asp Ser Ala Ser Val Pro 265 Glu Thr Pro Leu Asp Cys Glu Val Ser Leu Trp Ser Ser Trp Gly Leu 280 Cys Gly Gly His Cys Gly Arg Leu Gly Thr Lys Ser Arg Thr Arg Tyr Val Arg Val Gln Pro Ala Asn Asn Gly Ser Pro Cys Pro Glu Leu Glu 305 310 Glu Glu Ala Glu Cys Val Pro Asp Asn Cys Val 325 <210> 41 <211> 330 <212> PRT <213> Rattus norvegicus <400> 41 Met Glu Asn Val Ser Phe Ser Leu Asp Arg Thr Leu Trp Val Phe Leu

Leu Ala Met Leu Gly Ser Thr Ala Gly Gln Pro Leu Gly Gly Glu Ser

Val Cys Thr Ala Arg Pro Leu Ala Arg Tyr Ser Ile Thr Phe Thr Gly Lys Trp Ser Gln Thr Ala Phe Pro Lys Gln Tyr Pro Leu Phe Arg Pro Pro Ala Gln Trp Ser Ser Leu Leu Gly Ala Ala His Ser Ser Asp Tyr Ser Met Trp Arg Lys Asn Glu Tyr Val Ser Asn Gly Leu Arg Asp Phe Ala Glu Arq Gly Glu Ala Trp Ala Leu Met Lys Glu Ile Glu Ala Ala Gly Glu Lys Leu Gln Ser Val His Ala Val Phe Ser Ala Pro Ala Val 120 Pro Ser Gly Thr Gly Gln Thr Ser Ala Glu Leu Glu Val His Pro Arg 135 His Ser Leu Val Ser Phe Val Val Arg Ile Val Pro Ser Pro Asp Trp 150 Phe Val Gly Ile Asp Ser Leu Asp Leu Cys Glu Gly Gly Arg Trp Lys Glu Gln Val Val Leu Asp Leu Tyr Pro His Asp Ala Gly Thr Asp Ser 180 185 Gly Phe Thr Phe Ser Ser Pro Asn Phe Ala Thr Ile Pro Gln Asp Thr Val Thr Glu Ile Thr Ala Ser Ser Pro Ser His Pro Ala Asn Ser Phe 215 Tyr Tyr Pro Arg Leu Lys Ser Leu Pro Pro Ile Ala Lys Val Thr Phe 225 Val Arg Leu Arg Gln Ser Pro Arg Ala Phe Ala Pro Pro Ser Leu Asp 250 Leu Ala Ser Arg Gly Asn Glu Ile Val Asp Ser Leu Ser Val Pro Glu 265 260 Thr Pro Leu Asp Cys Glu Val Ser Leu Trp Ser Ser Trp Gly Leu Cys Gly Gly Pro Cys Gly Lys Leu Gly Ala Lys Ser Arg Thr Arg Tyr Val 295 Arg Val Gln Pro Ala Asn Asn Gly Thr Pro Cys Pro Glu Leu Glu Glu 315 320 305 Glu Ala Glu Cys Ala Pro Asp Asn Cys Val 325

- <210> 42
- <211> 331
- <212> PRT
- <213> Danio rerio
- <400> 42
- Met Glu Thr Met Thr Ser Leu Arg Val Asn Cys Trp Leu Thr Met Thr 1 5 10 15
- Leu Ala Leu Leu Ser Gly Val Pro Ala Met Pro Val Asp Val Asp Arg
  20 25 30
- Met Cys Thr Ala Pro Ser Thr Ala Lys Tyr Arg Leu Thr Phe Thr Gly 35 40 45
- Gln Trp Thr Gln Thr Ala Phe Pro Lys His Tyr Pro Leu Tyr Arg Pro 50 55 60
- Pro Ala Gln Trp Ser Pro Leu Ile Gly Val Thr His Ser Ser Asp Tyr
  65 70 75 80
- His Leu Trp Gln Arg Asn Glu Tyr Ala Ser Asn Gly Val Arg Glu Phe
  85 90 95
- Ser Glu Arg Ala Glu Ala Trp Thr Leu Ile Lys Glu Val Glu Ala Ala 100 105 110
- Gly Glu Arg Ile Gln Ser Val Tyr Gly Leu Phe Ser Ala Pro Ala Val 115 120 125
- Val Ala Gly Thr Gly His Ala Thr Thr Glu Phe Glu Val Phe Ala Arg 130 135 140
- His Ser Leu Leu Ser Phe Ile Val Arg Ile Val Pro Ser Pro Asp Trp 145 150 155 160
- Phe Ile Gly Val Asp Ser Leu Asn Leu Cys Glu Gly Asp His Gly Lys 165 170 175
- Glu Asn Ile Ser Leu Glu Leu Tyr Pro Tyr Asp Ala Gly Thr Asp Ser 180 185 190
- Gly Phe Thr Phe Ser Ser Pro Asn Phe Glu Thr Ile Pro Gln Asp Lys 195 200 205
- Val Thr Gln Ile Thr Ser Ser Phe Pro Ser His Pro Ala Asn Ser Phe 210 215 220
- Tyr Tyr Pro Arg Leu Lys His Leu Pro Pro Ile Ala Lys Val Ser Leu 225 230 235 240
- Thr Lys Ile Lys Asn Asn Gln Ile Phe Ser Leu Pro Ile Gln Pro Thr 245 250 255
- Gln Ser Asn Gln Ile Pro Ser Gly Asn Glu Ile Asp Gly Pro Leu Ile

Asn Thr Pro Leu Asp Cys Glu Val Ser Val Trp Ser Pro Trp Gly Leu 275 280 285

Cys Lys Gly Gln Cys Gly Glu Lys Gly Val Lys His Arg Thr Arg Tyr 290 295 300

Ile His Met His Pro Ala Asn Asn Gly Ala Pro Cys Pro Ser Leu Glu 305 310 315 320

Glu Lys Arg Leu Cys Ile Pro Asp Asn Cys Val 325 330

<210> 43

<211> 104

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Consensus Sequence

<400> 43

Pro Ser Phe Leu Val Ser Gly Thr Phe Asp Ala Arg Gly Gly Arg Leu 1 5 10 15

Arg Gly Pro Arg Thr Gly Val Arg Leu Ile Ile Pro Pro Gly Ala Ile 20 25 30

Pro Gln Gly Thr Arg Tyr Thr Cys Tyr Leu Val Val His Asp Lys Leu
35 40 45

Ser Thr Pro Pro Pro Leu Glu Glu Glu Glu Thr Leu Leu Ser Pro Val 50 60

Val Glu Cys Gly Pro His Gly Ala Leu Phe Leu Arg Pro Val Ile Leu 65 70 75 80

Glu Val Pro His Cys Ala Ser Leu Arg Pro Arg Asp Trp Glu Ile Val 85 90 95

Leu Leu Arg Ser Glu Asn Gly Gly
100

<210> 44

<211> 104

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: ZU5 domain Consensus Sequence

<400> 44

Ser Gly Phe Leu Val Ser Gly Thr Phe Asp Ala Arg Gly Gly Arg Leu
1 5 10 15

Arg Gly Pro Arg Thr Gly Val Arg Leu Ile Ile Pro Pro Gly Ala Ile 20 25 30

Pro Gln Gly Thr Arg Tyr Thr Cys Tyr Leu Val Val His Asp Lys Leu 35 40 45

Ser Thr Pro Pro Pro Leu Glu Glu Gly Glu Thr Leu Leu Ser Pro Val 50 55 60

Val Glu Cys Gly Pro His Gly Ala Leu Phe Leu Arg Pro Val Ile Leu 65 70 75 80

Glu Val Pro His Cys Ala Ser Leu Arg Pro Arg Asp Trp Glu Leu Val 85 90 95

Leu Leu Arg Ser Glu Asn Gly Gly
100

<210> 45

<211> 96

<212> PRT

<213> Artificial Sequence

<220>

<400> 45

Pro Pro Gly Ala Ala Ser Leu Thr Glu Leu Thr Arg Glu Lys Leu Ala
1 5 10 15

Lys Leu Leu Asp His Asp Leu Gly Asp Asp Trp Arg Glu Leu Ala Arg
20 25 30

Lys Leu Gly Leu Ser Glu Ala Asp Ile Asp Gln Ile Glu Thr Glu Ser 35 40 45

Pro Arg Asp Leu Ala Glu Gln Ser Tyr Gln Leu Leu Arg Leu Trp Glu 50 55 60

Gln Arg Glu Gly Lys Asn Ala Thr Leu Gly Thr Leu Leu Glu Ala Leu
65 70 75 80

Arg Lys Met Gly Arg Asp Asp Ala Val Glu Leu Leu Arg Ser Glu Leu 85 90 95

<210> 46

<211> 51

<212> PRT

```
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: TSP1 domain
      Consensus Sequence
<400> 46
Trp Gly Glu Trp Ser Glu Trp Ser Pro Cys Ser Val Thr Cys Gly Gly
Gly Val Gln Thr Arg Thr Arg Cys Cys Asn Pro Pro Pro Asn Gly Gly
                                  25
Gly Pro Cys Thr Gly Pro Asp Thr Glu Thr Arg Ala Cys Asn Glu Gln
                              40
Pro Cys Pro
     50
<210> 47
<211> 83
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Death domain
      Consensus Sequence
<400> 47
Arg Glu Leu Cys Lys Leu Leu Asp Asp Pro Leu Gly Arg Asp Trp Arg
                  5
                                      10
Arg Leu Ala Arg Lys Leu Gly Leu Ser Glu Glu Glu Ile Asp Gln Ile
Glu His Glu Asn Pro Arg Leu Ala Ser Pro Thr Tyr Gln Leu Leu Asp
Leu Trp Glu Gln Arg Gly Gly Lys Asn Ala Thr Val Gly Thr Leu Leu
     50
Glu Ala Leu Arg Lys Met Gly Arg Asp Asp Ala Val Glu Leu Leu Glu
                     70
                                          75
Ser Ala Leu
<210> 48
<211> 86
<212> PRT
<213> Artificial Sequence
```

186

<223> Description of Artificial Sequence: IG domain

Consensus Sequence

<220>

<400> 48

Pro Pro Ser Val Thr Val Lys Glu Gly Glu Ser Val Thr Leu Ser Cys
1 5 10 15

Glu Ala Ser Gly Asn Pro Pro Pro Thr Val Thr Trp Tyr Lys Gln Gly
20 25 30

Gly Lys Leu Leu Ala Glu Ser Gly Arg Phe Ser Val Ser Arg Ser Gly 35 40 45

Gly Asn Ser Thr Leu Thr Ile Ser Asn Val Thr Pro Glu Asp Ser Gly 50 55 60

Thr Tyr Thr Cys Ala Ala Thr Asn Ser Ser Gly Ser Ala Ser Ser Gly 65 70 75 80

Thr Thr Leu Thr Val Leu 85

<210> 49

<211> 51

<212> PRT

<213> Artificial Sequence

<2205

<223> Description of Artificial Sequence: LRRCT domain Consensus Sequence

<400> 49

Asn Pro Phe Ile Cys Asp Cys Glu Leu Arg Trp Leu Leu Arg Trp Leu 1 5 10 15

Arg Glu Pro Arg Arg Leu Glu Asp Pro Glu Asp Leu Arg Cys Ala Ser

Pro Glu Ser Leu Arg Gly Pro Leu Leu Glu Leu Leu Pro Ser Asp Phe 35 40 45

Ser Cys Pro 50

<210> 50

<211> 22

<212> DNA

<213> Artificial Sequence

-22A-

<400> 50

ctgcacttca aggacagtta cc

22

```
<210> 51
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Ag1395 Probe
       Primer
<400> 51
ctatccatcc acgatgtgcc cagct
                                                                     25
<210> 52
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Ag1395 Reverse
      Primer
<400> 52
tgacaaggag cttactcttc ca
                                                                     22
<210> 53
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Ag395 Forward
      Primer
<400> 53
caggaagaaa taagccaagt cca
                                                                    23
<210> 54
<211> 19
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Ag395 Probe
      Primer
<400> 54
tccttggcct cccgcctgc
                                                                    19
<210> 55
<211> 24
<212> DNA
<213> Artificial Sequence
```

```
<220>
<223> Description of Artificial Sequence: Ag395 Reverse
       Primer
<400> 55
gaggtcatgt tctagcttcc catt
                                                                     24
<210> 56
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Ag888 Forward
      Primer
<400> 56
catagctgac cgcatctgaa
                                                                     20
<210> 57
<211> 26
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Ag888 Probe
      Primer
<400> 57
aatgctccat ctccttggct gagtga
                                                                    26
<210> 58
<211> 21
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Ag888 Reverse
      Primer
<400> 58
ggagctagca tccatcatca c
                                                                    21
<210> 59
<211> 21
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Ag784 Forward
      Primer
<400> 59
```

gtcct	tgg	gat	gtgt	gaga	iga t	:										21
<211: <212:	10> 60 11> 26 12> DNA 13> Artificial Sequence															
<220 × <223 ×	23> Description of Artificial Sequence: Ag784 Probe Primer															
<400> 60 cagagagacg cagctcctcc aagaag														26		
<210 > <211 > <212 > <213 >	> 22 > Di	1 <b>A</b>	icia	l Se	quen	ce										
<220> <223>	· De	escr rime:		on o	f Ar	tifi	cial	Seq	uenc	e: A	g784	Rev	erse			
<400> gaaca			acaga	agct	tc a	g										22
<210><211><211><212><213>	10 PF	)4 ?T	icial	l Se	quen	ce										
<220> <223>	De		iptio nsus				cial	Seq	uence	e: Z	U5 do	omai	n			
<400> Pro S 1			Leu	Val 5	Ser	Gly	Thr	Phe	Asp 10	Ala	Arg	Gly	Gly	Arg 15	Leu	
Arg G	ly	Pro	Arg 20	Thr	Gly	Val	Arg	Leu 25	Ile	Ile	Pro	Pro	Gly 30	Ala	Ile	
Pro G	ln	Gly 35	Thr	Arg	Tyr	Thr	Cys 40	Tyr	Leu	Val	Val	His 45	Asp	Lys	Leu	
Ser T	hr 50	Pro	Pro	Pro	Leu	Glu 55	Glu	Gly	Glu	Thr	Leu 60	Leu	Ser	Pro	Val	
Val G	lu	Cys	Gly	Pro	His 70	Gly	Ala	Leu	Phe	Leu 75	Arg	Pro	Val	Ile	Leu 80	
Glu Va	al	Pro	His	Cys 85	Ala	Ser	Leu	Arg	Pro 90	Arg	Asp	Trp	Glu	Ile 95	Val	

Leu Leu Arg Ser Glu Asn Gly Gly 100